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DEVELOPMENT AND CHARACTERIZATION OF A TRIDIMENSIONAL INTESTINAL MODEL TO STUDY PROTEIN DRUGS ABSORPTION

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To engineering. To science.

ABSTRACT

Despite the huge palette of the existing intestinal models, the drug development industry is not entirely satisfied since these models poorly replicate the architecture and mechanisms occurring in a living tissue. The current understandings on a number of biological processes are based on two-dimensional (2D) surfaces however *in vivo* cells primarily exist in a heterogeneous and information-rich environment which modulate a diversity of cellular events. The 2D models are a reductionist approach that not faithfully captures the *in vivo* scenario. In fact, the striking similarity between the *in vivo* morphology and behaviour of cells in tridimensional (3D) culture conditions, have turn 3D models in a mainstream approach.

The present study extends a traditional intestinal model to the deeper layers of the small intestine, in particular the intestinal mucosa that includes a broad variety of stromal cells. Fundamentally, this work aimed to dissect the stromal-epithelial interactions and evaluate the permeation of a protein drug, insulin, using a 3D based model comprising CCD18-Co intestinal myofibroblasts embedded in Matrigel, Caco-2 and HT29-MTX epithelial cells and Raji B lymphocytes.

As the 3D environment triggers different cellular responses of the 2D environment, the density of CCD18-Co cells was optimized and the results showed that 1×10^4 cells/cm² was the cell density better adapted to 21 days of culture, period necessary to fully differentiate Caco-2 cells. CCD18-Co cells embedded in Matrigel showed to exert a supportive role in the maintenance of the epithelial cells onto them. This structure is similar the native intestinal mucosa architecture. Matrigel, primarily acting as an initial livelihood, was substituted by extracellular matrix components newly produced by CCD18-Co cells.

All the studied models revealed functionally effective, returning sustainable results of transepithelial electrical resistance and expression of tight junction (TJ) proteins, which are indicators of membrane integrity and tightness respectively. Although Caco-2 cells form tight junctions between them, the introduction of mucus-producing HT29-MTX cells interferes with the tightness of the barrier due to structural differences between the TJs.

As mean to explore the transcytotic capacity of M cells, Raji B lymphocytes were added to the basolateral side of the Transwell insert to carry out the phenotypic conversion of Caco-2 cells into M-like cells. Although trigger these cells is a challenge due to the inexistence of a distinct cell marker, the lack of microvilli was the best cue to identify M-like cells.

The 3D quadruple model showed to be efficient in the prediction of insulin permeability. The less tight junctions between the epithelial cells, the existence of mucus and the presence of M-like cells were critical factors in the transport of insulin.

Ultimately, this model closely resembles the intestinal mucosa representing a sophisticated and robust tool useful in the absorption's evaluation of incoming drugs in the market.

RESUMO

Apesar da existência de uma grande paleta de modelos intestinais, a indústria de desenvolvimento de novos fármacos não está inteiramente satisfeita, uma vez que estes modelos não replicam plenamente a arquitetura e os mecanismos existentes num organismo. A atual compreensão de muitos processos biológicos é baseada em superfícies bidimensionais (2D), no entanto, *in vivo*, as células estão rodeadas por um ambiente heterogêneo e informativo que modula uma diversidade de eventos celulares. Os modelos 2D são uma abordagem reducionista que não captura fielmente o cenário *in vivo*. Na verdade, a semelhança notável da morfologia e comportamento *in vivo* em condições de cultura tridimensional (3D) tem tornado os modelos 3D a principal abordagem.

O presente estudo estende um modelo intestinal tradicional para as camadas mais profundas do intestino delgado, em particular a mucosa intestinal, que inclui uma ampla variedade de células do estroma. Fundamentalmente, este trabalho visou dissecar as interações estroma-epitélio e avaliar a permeação de um fármaco proteico, a insulina, usando um modelo 3D composto por miofibroblastos intestinais CCD18-Co embebidos em Matrigel, células epiteliais Caco-2 e HT29-MTX e linfócitos Raji B.

Como o ambiente 3D desencadeia diferentes respostas celulares do ambiente 2D, a densidade de células CCD18-Co foi otimizada e os resultados demonstram que 1×10^4 células/cm² foi a densidade celular melhor adaptada aos 21 dias de cultura, período necessário para diferenciar totalmente as células Caco-2. As células CCD18-Co embebidas em Matrigel mostram exercer um papel de suporte na manutenção das células epiteliais. Esta estrutura é semelhante à arquitetura nativa da mucosa intestinal. Matrigel, que inicialmente atuava como um meio de vida, foi substituído por componentes da matriz extracelular recém produzidos pelas células CCD18-Co.

Todos os modelos estudados se revelaram funcionalmente eficazes, devolvendo resultados sustentáveis de resistência transepitelial e expressão de junções apertadas, que são indicadores de integridade e estreiteza da membrana, respetivamente. Apesar das células Caco-2 estabelecerem junções intercelulares entre elas, a introdução das células HT29-MTX produtoras de muco interfere com a estreiteza da barreira devido às diferenças estruturais existentes entre as junções.

Como meio de explorar a capacidade transcitótica das células M, linfócitos Raji B foram adicionadas na parte basolateral do Transwell insert para conduzir a conversão fenotípica das células Caco-2 em células do tipo M. Embora detetar estas células seja um desafio devido à inexistência de um marcador distinto, a falta de microvilosidades foi a melhor pista para identificar as células do tipo M.

O modelo quádruplo 3D mostrou ser eficiente na previsão da permeabilidade da insulina. O carácter menos apertado entre as células epiteliais, a existência de muco e a presença das células M foram fatores críticos no transporte de insulina.

Para finalizar, este modelo assemelha-se à mucosa intestinal, representando uma ferramenta sofisticada e robusta útil para a avaliação da absorção de novos fármacos que chegam ao mercado.

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LIST OF ABBREVIATIONS

2D - Two-dimensional

3D - Tridimensional

ADME - Administration, Distribution, Metabolism, Excretion

AJs - Adherent junctions

ALP - Alkaline phosphatase

AMPs - Antimicrobial proteins

APES - 3-aminopropyltriethoxysilane

BM - Basement membrane

CCL20 - Chemokine ligand 20

CCR6 - Chemokine receptor 6

CD - Cluster of differentiation

CYP450 - Cytochrome P450

DAPI - 4', 6-diamidino-2-phenylindole

DCs - Dendritic cells

DMEM - Dulbecco's Modified Eagle's Medium

DMSO - Dimethyl sulfoxide

EDTA - Ethylenediamine tetraacetic acid

FAE - Follicle-associated epithelium

FBS - Fetal Bovine Serum

FN - Fibronectin

GALT - Gut-associated lymphoid tissue

GI - Gastrointestinal

HBSS - Hanks Balanced Salt Solution

HDMS - Hexamethyldisilazane

HGF - Hepatocyte growth factor

IDE - Insulin-degrading enzyme

IEL - Intraepithelial lymphocytes

IESC - Intestinal epithelial stem cells

IL - Interleukin

IMFs - Intestinal myofibroblasts

JAMs - Junctional adhesion molecules

KGF - Keratinocyte growth factor

LPLs - Lamina propria lymphocytes

LT α 1 β 2 - Lymphotoxin

LT β R - Lymphotoxin- β receptor

M cells - Microfold cells

MALT - Mucosa-associated lymphoid tissue

MDCK - Madin Darby Canine Kidney

MDR - Multidrug resistance

M-like cells - Microfold-like cells

MLNs - Mesenteric lymph nodes

MMPs - Metalloproteinases

MRP - Multidrug resistance-associated proteins

MTX - Methotrexate

MUC - Mucin

NEAA - Non-essential aminoacids

PBS - Phosphate Buffer Solution

PDL - Poly-D-Lysine

P-gp - P-glycoprotein

PI - Propidium Iodide

PPs - Peyer's Patches

PrP^c - Cellular prior protein

RANK - Receptor activator of nuclear factor kappa-B

RANKL - Receptor activator of nuclear factor kappa-B ligand

RT - Room temperature

SED - Subepithelial dome

SEM - Scanning electron microscopy

SF⁴ - Scatter growth factor

SLAA - Sialylated Lewis A antigen

TE - Tris-EDTA

TEER - Transepithelial electrical resistance

TEM - Transmission electron microscopy

TFF - Trefoil factors

TJs - Tight junctions

TNF- α - Tumor necrosis factor - α

UMOD - Uromodulin

ZO - Zonula occludens

α -SMA - α -smooth muscle actin

CHAPTER I

INTRODUCTION

1.1. Intestinal and mucosal functioning

Intestine is the major portion of gastrointestinal (GI) tract whose primary function is to gather the entrance of pathogens, toxins and undigested molecules while enabling the digestion and selective absorption of essential nutrients into the body. These distinct functions place the intestinal mucosa in the center of interactions between the immune system and the luminal contents [1]. There is no doubt that there is a tight control of enteric bacteria population and the luminal contents presenting those components to diverse physical and chemical intestinal barriers. If the foremost barriers fail, immune cells (phagocytes and lymphocytes underlying the *lamina propria*) are the next in line to carry out defensive actions [2].

Human small intestine, owing the larger surface area provided by valves of Kerckring (macroscopic valve-like folds), villi (finger-like protrusions), crypts (finger-like invaginations) and microvilli (cytoplasmatic extension), represents the protagonist organ in the drug absorption [3]. Absorption evaluation plays an ever increasingly role at the early stages of drug discovery due to its colossal potential to eliminate molecules with poor ADME (Absorption, Distribution, Metabolism and Excretion) properties from the drug development pipeline. Therefore, the replication of the intestinal functioning has revealed an attractive strategy that will potentially energize the drug development.

1.2. Intestinal progeny

The intestinal epithelium is unique in the cell proliferation, differentiation and apoptosis mechanisms as it occurs in an orderly fashion along the crypt-villus axis but ceases as cells reach the apex of villi ongoing spontaneous apoptosis and detaching into the lumen [4]. Wnt signal transduction pathway is the primary driving force behind intestinal proliferation responsible for maintain the crypt cell population in proliferative state [5, 6]. Proliferative crypts are monoclonal compartments [7] that harbour intestinal epithelial stem cells (IESC) and their progeny - transit-amplifying cells - partially differentiated cells that divide 4-5 times before terminally differentiate (**Fig. 1.1 A**). The differentiated cells generated from this process are divided in two populations that holds absorptive enterocytes and secretory cells including goblet, enteroendocrine and Paneth cells (**Fig. 1.1 B**) [5]. Although most reviews are focused on the aforementioned four cell types, there are described three additional intestinal cell types including microfold cells (M cells) [8], tuft cells or brush cells [9] and cup cells [10].

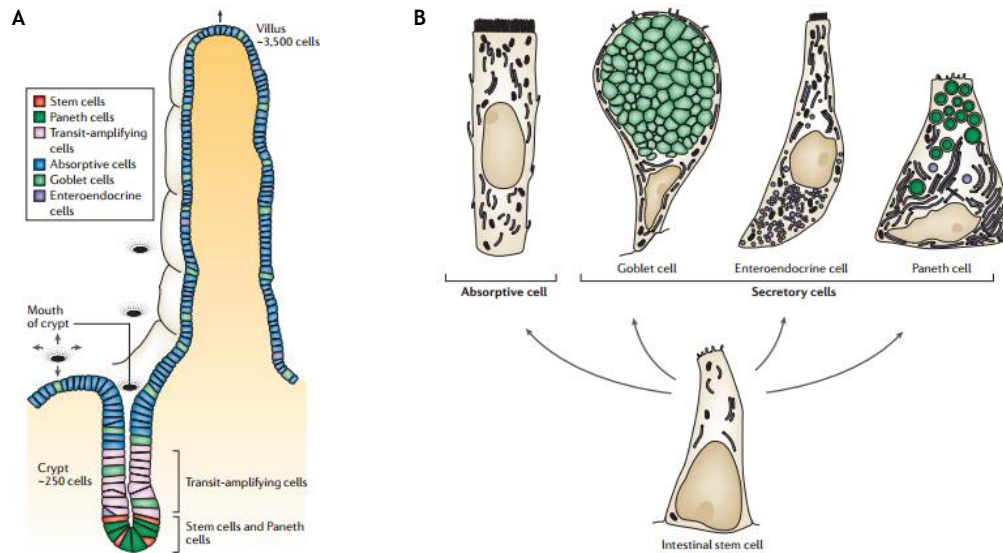


Figure 1. 1. Intestinal cells lineage in mammalian small intestine. Migratory process of transit-amplifying cells occurs in an orderly fashion along the crypt-villus (A) given origin to four main cell types classified according absorptive or secretory cells (B). Adapted from Crosnier C. et al. [4].

Absorptive enterocytes, representing about 80% of intestinal epithelial cells [11], are columnar and polarized cells discernible by the presence of apical brush border relevant in nutrient absorption. Goblet cells, scattered throughout the epithelium, are responsible for the secretion of protective mucins and trefoil factors (TFF) required for the movement and expulsion of gut contents and provide protection against shear stress and chemical damage. Enteroendocrine cells, about ~1% of epithelial cell population, represent a bridge between the central and enteric neuroendocrine system throughout the secretion of peptide hormones [12]. Paneth cells, with a function in innate immunity [13], are the only differentiated cells that reside on the undifferentiated compartment of intestinal mucosa, the crypts. These cells are usually identified by the large apical secretory granules that contain lysozymes, antimicrobials proteins (AMPs) and defensins [14]. Collectively, the diverse functions of intestinal epithelial cells result in a dynamic barrier that maintains the segregation between the luminal microbial communities and the mucosal immune system (Fig. 1.2) [12].

Overall, intestinal stem cells, their progeny and their microenvironment make up the niche, an anatomical and functional structure that coordinates the normal homeostatic production of functional mature cells. An intimate regulation between intrinsic and extrinsic cellular mechanisms maintains the balance between self-renewal and differentiation events [7].

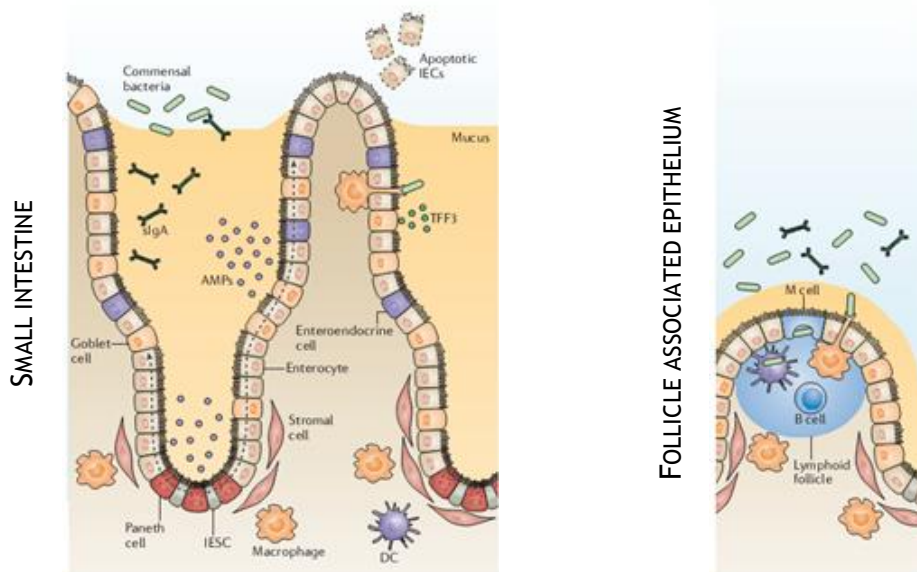


Figure 1. 2. The regulatory barrier function of intestinal epithelial cells. AMPs and TFF, respectively secreted by goblet and Paneth cells, promote the exclusion of bacteria from the epithelial surface. The luminal release of secretory IgA (sIgA) protects the intestinal epithelium from enteric toxins and pathogenic microorganisms. M cells mediate the transport of luminal antigens and bacteria across the epithelial barrier for immunological surveillance. Adapted from Peterson L. and Artis D. [12].

1.3. Intestinal immunology

Despite the selective character provided for highly efficient barriers, some pathogens can overcome the protective mechanisms and enter in the systemic circulation. As an extra-protective effort, the intestinal lining has more lymphoid cells and produces more antibodies than any other organ in the body [15]. The complex interplay of regulatory systems ensures the maintenance of gut homeostasis allowing the clear discrimination between invasive organisms and harmless antigens. Indeed, an active immunity against non-pathogenic materials and hypersensitivity against dietary antigens and commensal bacteria can unroll in inflammatory disorders such as coeliac disease and Crohn's disease, respectively [1].

Every mucosa region in the body has their own region for immunological surveillance known as mucosa-associated lymphoid tissue (MALT), with similar but not identical cellular composition [16]. The intestinal immune responses are entrusted to gut-associated lymphoid tissue (GALT), which tasks are divided for induction sites consisting of PPs and mesenteric lymph nodes (MLNs), involved in the induction phase of the immune response [2, 17] and effector sites that are diffused throughout the lamina propria and the epithelium of the mucosa [15]. Both the PPs and the villus are drained by afferent lymphatic that go to MLNs [1] (Fig. 1.3).

PPs have been increasingly valued since they are considered the immune sensors of intestine [18]. They are macroscopic lymphoid aggregates found in submucosa surrounded by a single layer of columnar epithelial cells, the follicle-associated epithelium (FAE), an interface between GALT and the luminal environment [18]. Mature PPs generally consists of

collections of large B-cell follicles and intervening T-cell areas [1]. Added to that, FAE is considered a selective break of intestinal barriers due to the lower levels of digestive enzymes, a less pronounced brush border surface and mucus production in reduced quantities, facilitating the access to the apical surface [18].

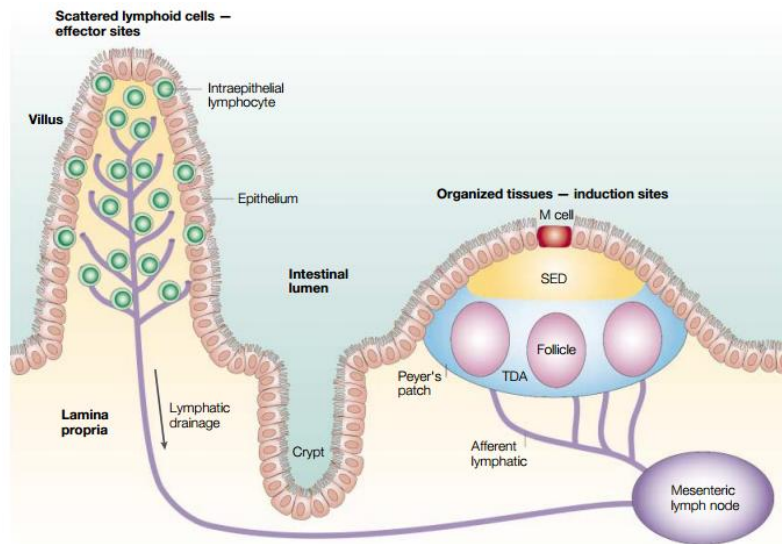


Figure 1. 3. Intestinal sites of immune response. The lymphoid areas are separated from the intestinal lumen by FAE and a more diffuse area below the epithelium known as subepithelial dome (SED). Adapted from Mowat A. [1].

1.3.1. M cells associated to lymphoid tissues

FAE's most determinant feature is the presence of M cells, specialized enterocytes for antigen sampling that constitute 10% of epithelial cells of the FAE [19]. However they represent a ratio of 1 in 10 million to the total number of intestinal epithelial cells [20]. These cells are highly specialized for the phagocytosis and transcytosis of luminal content, particularly antigens and pathogenic or commensal microorganisms [21, 22]. Beneath the M-cell basolateral membrane, antigens are taken up by intraepithelial pocket which contains a population of lymphocytes and mononuclear phagocytes (a heterogeneous population of macrophages and classical DCs) (**Fig. 1.4 A**) [8].

Morphologically distinct from enterocytes, M cells display poorly organized brush border with short irregular microvilli at the apical domain (**Fig. 1.4 B**), usually identified by immunostaining of actin and villin [23]. The usual thick glycocalyx of absorptive cells is replaced in M cells by a thin glycocalyx. Naturally, these features lead to a highly attenuated expression of apical digestive enzymes such as alkaline-phosphatase and iso-sucrase maltase [20, 24].

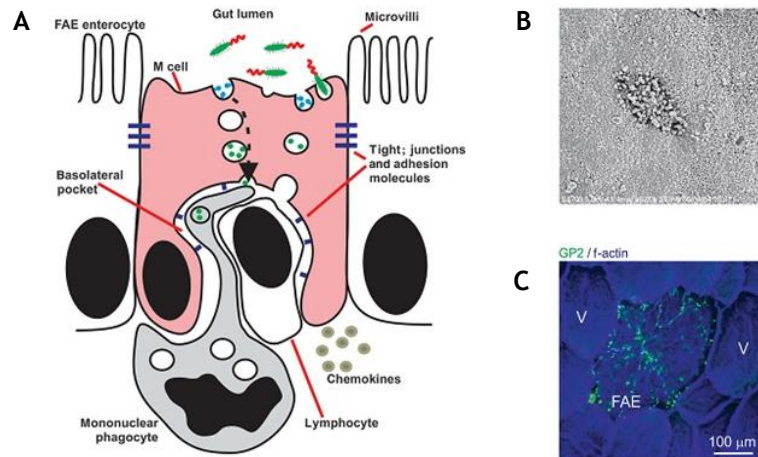


Figure 1.4. Morphological features of M cells. (A) M cell features denoting the lack in microvilli and mucus layer and the basolateral pocket containing immunosurveillance cells. (B) Scanning electron microscopy of M cell highlighting the blunted microvilli. (C) Immunohistochemical analysis of mature M cells in the FAE of a mouse PPs of specifically expressing glycoprotein 2 (GP2). Adapted from Mabbott N. A. *et al.* [8].

The reduced brush border and lack in enzymatic activity clarifies the inexistence of a prominent role of M cells in digestion or absorption. On the other hand, the exposed nature of M cell apical surface and their location sited over lymphoid follicles assuring the immediate exposure of foreign matter to underlying immune cells, points the transepithelial transport as the primary function of M cells [24].

Due to the lack of knowledge, general agreement on M cell function fails to go further than this, but, at least, M cells express a variety of “immunosurveillance” receptors on their apical surfaces that enable them to sample diverse microbial pathogens and antigens such as glycoprotein 2 (GP2), uromodulin (UMOD) and cellular prior protein (PrP^C), among others [8, 19, 25].

1.3.1.1. M cell differentiation in the gut epithelium

In the intestine, the dome-associated crypts surrounding the FAE and the villous crypts at the base of the villi contain cycling Lgr5⁺ cells (leucine-rich repeat-containing G protein couple receptor) intermingled among Paneth cells, from each almost epithelial cells lineage, including M cells, emerged (Fig. 1.5, A) [26]. As both dome associated crypts and villous crypts contain cycling Lgr5⁺ stem cells, additional signals from the stromal and immune cells in the SED are required to induce M cell differentiation. In intestine, the cytokine Receptor Activator of NF- κ B Ligand (RANKL) is selectively expressed by the subepithelial stromal cells beneath the FAE (Fig. 1.5, B) [8]. Analysis of a RANKL-deficient mice and administration of exogenous RANKL *in vivo* reveals that this cytokine is a critical factor that controls the differentiation of RANK-expressing enterocytes into M cells. Indeed, the interaction between RANKL and its receptor RANK will induce Spi-B (transcription factor) expression that could induce the maturation of M cells (Fig. 1.5, C) [27, 28].

M cell phenotype can also be induced by immune cells attracted to the dome region by the cytokine CCL20 produced by epithelial cells (Fig. 1.5, D) [29]. Immune cells interact with CCL20 through their receptor CCR6, verifying that when this signalling is impeded, M cell maturation is likewise compromised [30]. However, CCL20-CCR6 signalling does not influence the initial M cell differentiation but instead hampers their functional maturation [31]. Therefore, the remaining question of how B cell interacts with M cells precursors is not fully clarified but some authors claim that CD37L-CD37 signalling between M cells progenitors and the basolateral pocket of B cells may provide at least one of the signals that induce final functional maturation [32].

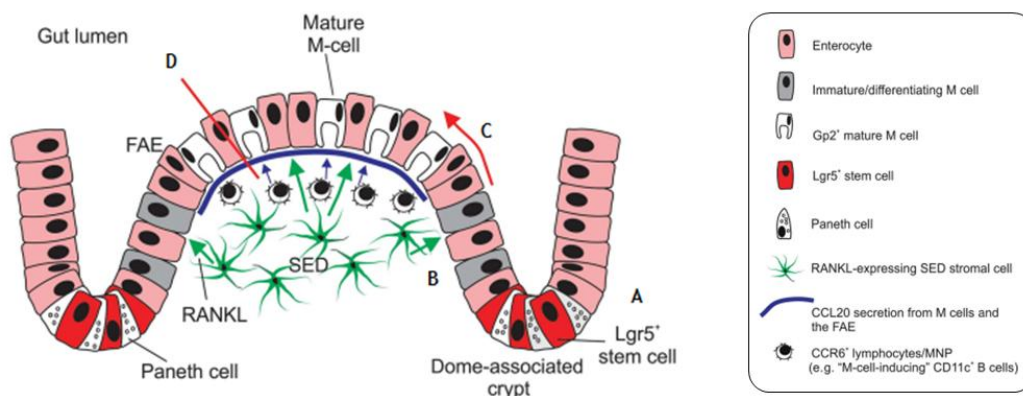


Figure 1. 5. Possible mechanism behind M-cell genesis derived from Lgr5⁺ stem cells in the crypts. (A) All the epithelial cell lineages develop from Lgr5⁺ intestinal stem cells located in the crypts. (B) RANKL is selectively expressed by the subepithelial stromal cells beneath the FAE. (C) The RANKL-RANK interaction will induce Spi-B expression that could induce the maturation of functional M cells. (D) Immune cells can also induce M cell phenotype attracted to the dome region by the cytokine CCL20 produced by epithelial cells. Adapted from Mabbott N. A. et al. [8].

1.3.1.2. *In vitro* M cell differentiation

The FAE that overlies Peyer's Patches constitutively expresses the chemokine CCL20 that is not expressed in the absorptive epithelium [33]. However the mechanisms behind the interplay between FAE differentiation and CCL20 expression are poorly understood. At the same time, lymphotoxin β receptor (LT β R) signalling plays a central role in the PPs' organogenesis [34, 35]. Taking together these events, *Rumbo et al.* studied whether LT β R mediates the CCL20 gene expression in intestinal epithelial cells. The results demonstrate *in vivo* and *in vitro* evidences that LT β R signalling may be responsible for the CCL20 production [33].

Other *in vitro* studies showed that lymphocytes trigger the conversion of enterocytes cell lines into M-like cells in co-culture under influence of membrane-bound lymphotoxin (LT α 182) secreted by B-lymphocytes. However, *in vivo* M cell differentiation reveals that the absence of mature lymphocytes does not prevent the formation of M cells indicating that signalling molecules that support M cell differentiation, such as lymphotoxin, may be supplied

by other underlying cells. Notwithstanding, mature lymphocytes are required for the formation of full-sized FAE [36].

It is still controversial whether M cells arise as a separate cell line or differentiate from enterocytes on the dome under the influence of immune cells [37, 38].

1.4. Epithelial - mesenchymal crosstalk

The function of the intestinal stem cells is to maintain the integrity of the intestinal epithelium. Stem cells dynamics are regulated by environmental factors, the sum of which constitutes the protective niche composed by neighbouring epithelial cells, mesenchymal cells, basement membrane (BM) and soluble cell- or matrix-associated growth factors [39]. It is established that although separated from the epithelial cells by the basement membrane, the constellation of mesenchymal elements promote the epithelial-mesenchymal crosstalk required to maintenance of the physiological homeostasis and the stem cell niche [40].

During the past decade, the permissive and instructive actions of the mesenchyme in the epithelial cells have been highlighted by the expression of important growth factors whose receptors are found on the epithelial cell surface, for instance, HGF/SF⁴ (hepatocyte growth factor/scatter factor) and KGF (keratinocyte growth factor) which regulate the epithelial cell behaviour [41].

1.4.1. Epithelial - stromal interface

As in most of the tissues, intestinal epithelium is supported by BM that separates epithelial cells from subepithelial fibroblasts or intestinal myofibroblasts (IMFs). IMFs are defined by phenotypic characteristics of both fibroblasts and smooth muscle cells [42]. They express α -smooth muscle actin (α -SMA), intermediate filaments such as vimentin but very weakly positive or negative for desmin and the absence of epithelial cytokeratins [43, 44] (Table 1.1). Although various combinations of these markers have been used to demonstrate the plasticity of IMFs, they are defined by their location and structure together with the expression of α -SMA, the best single but not absolutely exclusive marker [39].

Table 1. 1. Mesenchymal stromal cell markers. Adapted from Pinchuk I. V. et al. [43].

CELLS MARKERS	Myofibroblasts	Fibroblasts	Pericytes	<i>Muscularis mucosae</i>
α -SMA	+	-	+	+
Desmin	-	-	\pm	+
Vimentin	+	+	+	-
CD90	+	+	+	\pm

IMFs have flattened cell bodies, constituting a cellular network that ensheaths the lamina propria of the intestinal villi. In the small intestinal villus, α -SMA staining of subepithelial cells is less than those surrounding the crypts, and may even be lost all together in the mid- and upper villus [45]. Subepithelial cells stain weakly for desmin, suggesting the existence of α -SMA cells in the intestinal villus such as pericytes (mural cells that surround the capillaries of the lamina propria) [46]. Originally believed to be restricted to a two-dimensional (2D) network along the villi, IMFs are actually connected to α -smooth negative fibroblasts-like cells and pericytes forming a tridimensional (3D) network [47, 48] (Fig. 1. 6).

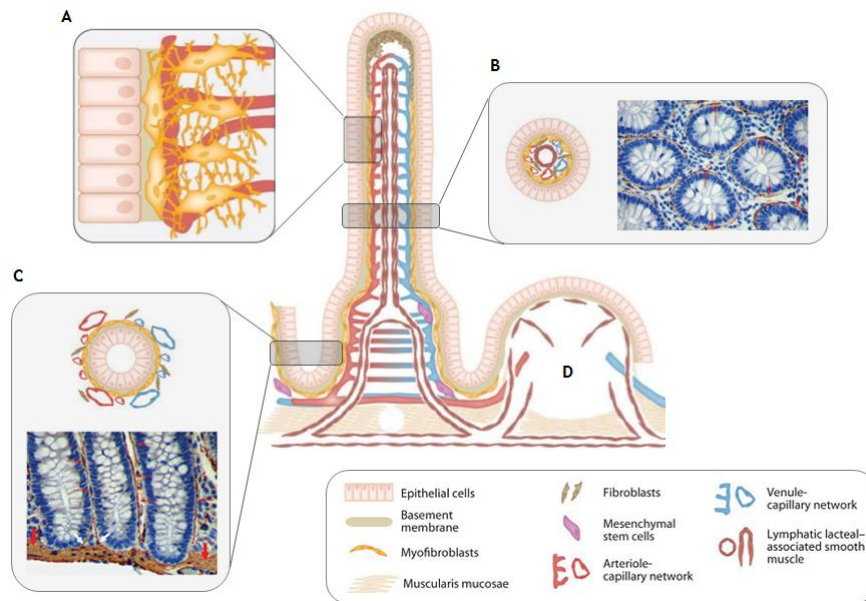


Figure 1. 6. Stromal-epithelial interactions along the villus-crypt axis. (A) Detailed interactions between IMFs that ensheaths the lamina propria and pericytes that support the capillaries. (B) Cross section through the intestinal villus with positive (brown) α -SMA IMFs. (C) Cross section through the crypts demonstrating the IMFs connected to the muscularis mucosae (thick red arrows) at the bases of the crypts. (D) PPs with its vascular, lymphatic and stromal elements, except lymphocytes, macrophages and dendritic cells. Adapted from Powell D. W. et al. [39, 42].

Several evidences state that the interaction between myofibroblasts and epithelial cells reveals truly important in supporting mucosal architecture and development but also the stem cell niche. IMFs have been associated to stem cell regulation proven to be necessary to the successful engraftment and proliferation of enteroids (intestinal crypt cultures) [49]. Although, these regulation is not clear due to the multiple factors and cell types known to be involved in stem cell regulation.

Additionally, IMFs are likely to play supportive roles promoting epithelial regrowth and enhancement of barrier function during time of injury throughout subepithelial myofibroblasts migration [50]. Likewise, the absence of myofibroblasts precluded the successful maintenance of epithelial cell formation and proliferation, even under the presence of supportive growth factors [51]. Tissue repair-associated factors such as $\text{TNF-}\alpha$, IL-1, IL-2, IL-6 and IL-15 activate myofibroblasts causing their motility and release of ECM components and growth factors. On the other hand, pro-inflammatory cytokines and growth factors control the expression of

matrix metalloproteinase (MMPs) and TIMP (specific inhibitors of MMPs) contributing to the tissue remodelling [48].

1.5. Intestinal barriers

The magnitude of interactions with the external environment reaches a pinnacle in the GI tract, the largest mucosal surface in continuous contact with dietary antigens and microorganisms. Before the immunological surveillance by the underlying immune cells, foreign organisms are exposed to protective barriers that will counteract absorptive events [52] (Fig. 1. 7).

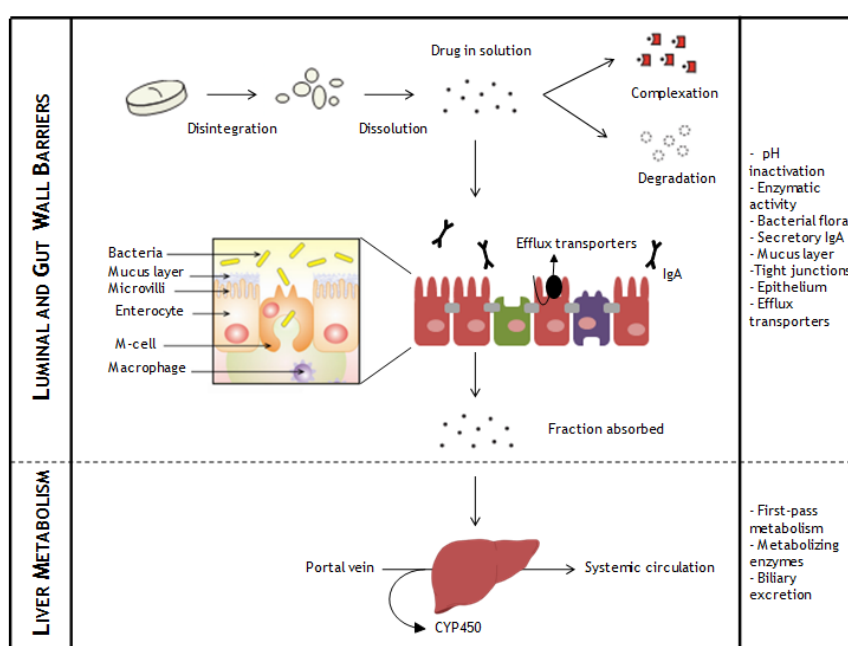


Figure 1. 7. Sequential barriers during the transit of a drug molecule in the GI tract when orally administered until reaches the systemic circulation. Adapted from Chen M. *et al.* and Artursson P. *et al.* [53, 54].

1.5.1. Luminal barriers

The drastic pH changes in stomach from slightly acid (pH 1.2 - 3.0) to slightly basic (pH 6.5 - 8.0) in the proximal end of ileum impact the drug stability and solubility [15]. Beyond small intestine being the major place of absorption, the enzymatic activity gathers a big palette of proteolytic enzymes divided in three classes: luminal, membrane-bound and cytosolic proteases. Oral administered compounds undergo proteolytic degradation in stomach by pepsin [55]. Upon reaching the duodenum, luminal proteases secreted by pancreas and released in the small intestine, namely serine endopeptidase, trypsin, α -chymotrypsin, elastase and exopeptidases (carboxypeptidases A and B), degrade the mixture of peptides and proteins. They are digested further by membrane-bound (brush-border) and cytosolic proteases (cytoplasm and lysosomes) associated to enterocytes. Proteases in the

brush border of enterocytes are potential the most biologically active enzymatic barrier, accounting with membrane-bound proteases and pancreatic enzymes that may be adsorbed from the luminal fluid [55, 56].

Apart from that, the aminopeptidases in PPs are about 20-30% of the neighbouring areas reinforcing that the unprotected FAE of PPs is potential targeting site for proteins delivery [57].

As the most abundant class of antibodies found in the intestinal epithelium, secretory IgA (sIgA) has been recognized as a first line defence in protecting the intestinal epithelium from enteric pathogens and toxins [58]. The production of sIgA is dependent on the sampling by M cells from PPs processing a pallet of antigen-presenting cells and multiple cytokines. sIgA operates in the clearance of antigens and pathogenic microorganisms from the intestinal lumen by blocking their access to epithelial receptors, a process known immune exclusion [59].

1.5.2. Gut wall barriers

Beyond the luminal barriers, the gut wall barriers are equally challenging consisting of mucus layer, the epithelium equipped with tight junctions and efflux transporters and metabolizing enzymes.

The entire intestine is covered by a stagnant layer composed by water, mucus and glycocalyx adjacent to the intestinal wall that is permeable to nutrients, water and small molecules but restrict to bacteria and pathogens [60]. Mucus secreted by the epithelial cells throughout the gastrointestinal tract has a unique secretion that forms a gel adherent to the surface constituting a protective wall. However, mucus layer is responsible by a rapid particle clearance due to the quick turnover, approximately 2-5 days in the small intestine of humans, preventing drug penetration and consequent absorption [61]. Additionally, the repulsion between drugs and the negatively charged mucus obstruct the direct contact between those drugs and the epithelial cells [62]. Even though all the limitations that mucus represents, its hydrogel-like structure reduces the shear stress and can stick a compound in the firmly adherent mucus layer increasing the transcellular migration into the epithelial wall [63].

The second rate-limiting step to absorption is the epithelium reinforced by tight junction and efflux transporters. Epithelial barrier is regulated by junctional complexes referred as tight junctions (TJs), adherent junctions (AJs), desmosomes and gap junctions. TJs between the neighbouring cells exclude the influx bacteria and molecules larger than 200 Da through paracellular routes [53] together with efflux transporters that forward unwanted substances back to the lumen. P-glycoprotein (P-gp) is the most studied member of multidrug resistance (MDR) family, primarily expressed on the apical membrane of the epithelial cells [64]. It reduces the intracellular accumulation or transcellular translocation of many compounds [65]. The combined role of P-glycoprotein with cytochrome P450 (CYP) 3A drug

metabolizing enzyme has been recognized as a major determinant of gut wall absorption [66, 67].

Finally, even if a significant amount of drug is transported through the intestinal epithelium, the first-pass metabolism by liver will highly reduce the dose fraction that reaches the systemic circulation [68]. Other limiting factors should fall in the scope of considerations including fast elimination from the systemic circulation, inducement of an immune response, uptake by non-target tissues and an inefficient target cell entry [60].

To overcome these barriers, innovative strategies have been proposed including chemical modification, formulation vehicles, proteases inhibitors, absorption enhancers and mucus-adhesive polymers [55, 60, 63].

1.6. Intestinal transport mechanisms

The process of absorption involves a set of processes that consists of pre-absorption, uptake and translocation and arrival to the destination targets. Molecules cross the intestinal epithelium having at disposal five distinct mechanisms: paracellular transport (A), transcellular passive transport (B), transcellular active transport or carrier mediated transport (C), receptor-mediated transcytosis (D) and via M cells of PPs (E), represented in Fig. 1. 8 [69].

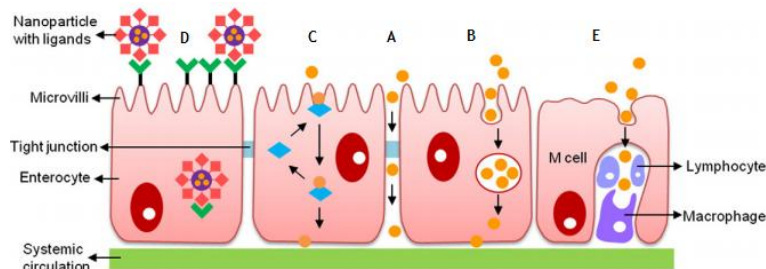


Figure 1. 8. Schematic representation of mechanisms of transport. (A) Paracellular transport; (B) Transcellular transport; (C) Carrier-mediated transport; (D) Receptor-mediated transport; (E) M-cell mediated transport by phagocytosis. Adapted from Yun et al. [55].

1.6.1. Paracellular pathway

Starting by the paracellular route, small hydrophilic particles may elect this route or the aqueous route that normally absorbs nutrients or vitamins [15], using the intracellular spaces between the epithelial cells (Fig. 1.8 A). This route is used by hydrophobic molecules such as oxygen and carbon dioxide that are small or neutral in charge and can cross the epithelial layer by uninhibited diffusion [70].

At a certain level, this route gained interest for peptide delivery due to the lack in proteolytic activity [60] however, the paracellular space represents less than 1% of the intestinal mucosa surface contributing for a minimal transport [70].

1.6.2. Transcellular pathway

Larger compounds are not able to use intracellular spaces, taking the transcellular route passively. Transcytosis or transcellular pathway involves the uptake in membrane-bound vesicles by a process named endocytosis at the apical membrane, further released in the basolateral membrane (**Fig. 1. 8 B**) [71]. The endocytic process is divided in phagocytosis (uptake of large particles) and pinocytosis (uptake of the fluids and solutes), the last divided at least in four basic mechanisms: macropinocytosis, clathrin-mediated endocytosis, caveolin-mediated endocytosis and clathrin and caveolin-independent endocytosis [72].

Notwithstanding, most of the therapeutic peptides and proteins are intact absorbed and are not expected to follow the transcellular route in a passive way. The active transcellular transport of peptides and proteins occurs via a specific carrier (**Fig. 1. 8 C**) [69]. Carrier-mediated pathway is a energy-dependent and saturable mechanism (concentration dependent) [73]. The molecule approaches the cell membrane and interacts with the transport proteins on the surface of the lipid bilayer forming a reversibly bound. These complex cross the lipid bilayer and when reaches the other side of the cell, the bound between the molecule and the transport protein is disrupted placing the molecule outside the cell [70].

A select group of macromolecules that are too large to slip through facilitated transport pathways and too constrained thermodynamically to transverse a lipid bilayer possible, readily moves across the transcellular barrier through receptor-mediated endocytosis (**Fig. 8 D**) [74]. This transport has been explored to increase the oral bioavailability of protein drugs by modification of receptor specific ligands with peptide and protein drugs. Protein drugs can act either as receptor specific ligands for surface-attached receptors or as receptor for surface-attached ligands [55]. This transport entails binding of the ligand to a specific cell surface receptor, receptor clustering and internalization [15].

1.6.3. M-cell mediated transport

As previously mentioned, the break in the intestinal barrier of the specialized epithelium FAE have been explored as a route of transport (**Fig. 1. 8 E**) [75]. M cells delivery by PPs involves the contact with the projections on M cells surfaces followed by a rapid phagocytosis. The fusion of the phagocytic vesicle with the basolateral membrane results in the emptying of the vesicle into the intraepithelial pocket. Particles can be engulfed by phagocytic cells in the pocket or pass through the basal lamina into the sub-epithelial region. Unavoidably, M cells mediated absorption represents an outstanding opportunity for explore drug absorption using nanocarriers [76].

1.7. Predicting intestinal absorption

Understanding and anticipating the mechanisms and factors that affect the gastrointestinal absorption is of utmost importance in the development and classification of new drugs [77].

With view to simplify and speed up the prediction of intestinal permeability, multiple screening techniques have been purposed. For many reasons, but particularly due to financial and ethical considerations, *in vitro* models are the most extensively used over the years based on Caco-2 intestinal epithelial cells [78]. Inherently, *in vitro* models raise questions about the extrapolation of results to the *in vivo* scenario. On the other hand, the complex *in vivo* environment difficult data interpretation together with the fact that, the microenvironment of a murine has subtle but not no negligible differences compared with *in vivo* circumstances [79].

What about artificial membranes and computational-based models? Are they on the way to replace Caco-2 models? In the drug development field, *in silico*, computational and mathematical models significantly support decisions about drug absorption in the early screening phase of drug evaluation. This power has a huge impact in drug companies, reducing the time and the costs in selecting potential candidates and performing the biological screening. The efficiency of this approach is recognized by the pharmaceutical industry however the insufficient database with reliable information is the major impediment. Therefore, the fact that neither the absorption process nor the experimental techniques used for data capture are not fully understood, the computational models will not do the job better [80, 81].

In this subchapter will be highlighted multiple approaches for drug absorption screening that reflect the increasingly necessity of predict the intestinal absorption, summarized in Table 1. 4 at the end of the subchapter.

1.7.1. *In vivo* models

No matter how sophisticated *in vitro* models are, *in vivo* evaluation will continue to be an invaluable tool required to validate the true performance of a drug or delivery system [82].

The main advantage of *in vivo* models is the integration of dynamic components such as the mesenteric blood circulation, the mucus layer and other factors that influence drug bioavailability [77].

In turn, the main weakness of *in vivo* models is the difficulty in separate the variables involved in the process of absorption, which difficult the identification of individual rate-limiting steps. The impact of anatomical and physiological differences, between and within species, influences the choice of the animal model [83]. For instance, the intestinal mucosal surfaces are comparable between species but drug transported proteins, enzymes and

microorganisms can differ. Rat is the most frequently animal model with similar absorption profiles and similar transport expression patterns to the human. However, the distinct patterns of metabolizing enzymes mistake the predictions [84]. The evolutionary proximity between monkeys and humans put this specie in a special place in drug discovery, however the costs and ethical concerns limit their use [77]. Pigs are also used to absorption studies due to the similarity of large intestine with human but data is rare [85]. Given these facts, has been stated that “the only real model for a man is man” [79].

1.7.2. *In vitro* models

It is been stated that *in vitro* models do not mimic the real situation, but these models cannot be reduced just to that. Cell-based models are a commitment between an intermediate level of complexity, with multiple transport systems than reflect the organ *in vivo* and what is feasible to reproduce in the laboratory [77]. In the field of drug discovery and development, what may be more relevant is to reproduce the extent and nature of transport and metabolising systems observed in the organ *in situ*, rather than the organ specific phenotype. In fact, animal models will always be part of drug development, clearly showing the propensity for oral absorption and the limitations that the whole organism offers to the absorption of a drug. At this point, *in vitro* models can sponsor a significant reduction of the *in vivo* models usage, allowing the earlier identification of drug candidates with poor ADME properties.

Therefore, the key thought is to understand the models limitations and use them as a mean to reduce the wastage and the expense of drug administered quantities, as well as to allow a more strategic planning of *in vivo* animal experiments.

1.7.3. Caco-2 based models

Since their isolation from a human colorectal carcinoma [86], Caco-2 cells have been widely used to absorption studies, considered to be the ‘gold standard’ of intestinal absorption assays [87]. Caco-2 cells spontaneously undergo typical enterocytic differentiation, a process that is growth related. At early stages of culture (3-4 days), Caco-2 cells maintained undifferentiated, displaying only few apical microvilli and do not express sucrase immunoreactivity (indicator of microvilli). At confluence they form polarized monolayers joined by tight junctions and presenting well-developed apical microvilli [88]. Despite their colonic origin, Caco-2 cells express the majority of morphological and functional characteristics of absorptive intestinal cells (Table 1. 2), inclusive enzymes such as disaccharidases and peptidases typically expressed by enterocytes. Adding to that, Caco-2 cells do not constitute a passive membrane, confirmed by identification of multiple uptake and efflux transporters [89]. However, the most important metabolizing enzyme in the human

gut, CYP3A4, is low or absent in Caco-2 cells, overestimating the fraction absorbed for substrates to 3A4 [90].

Some complications have been reported with respect to this cell line such as the high and variable expression of P-gp and low expression of metabolising enzymes. Added to that, the pore size of Caco-2 cells is estimated to be approximately 3.7 ± 0.1 Å, tremendously underestimating the paracellular transport [91]. The relatively wide variation of the correlation profiles between different laboratories and within the same laboratory appears to be a disadvantage, resultant of different culture conditions, protocols or genetic drift [81, 92].

Table 1. 2. Morphological and functional characteristics of Caco-2 cell line [79, 88, 90, 93].

MORPHOLOGY	Grow in monolayer, polarized morphology, microvilli on the apical side, tight junctions between the adjacent cells and express small intestinal hydrolase enzyme activities (sucrase-isomaltase, lactase, aminopeptidase N, dipeptidylpeptidase IV).
INTEGRITY PARAMETERS	Transepithelial electrical resistance after confluence and permeability marker compounds (mannitol).
MEMBRANE IONIC TRANSPORT	Na ⁺ /K ⁺ ATPase, H ⁺ /K ⁺ ATPase, Na ⁺ /H ⁺ exchange, Na ⁺ /K ⁺ /Cl ⁻ co-transport, apical Cl ⁻ channels.
MEMBRANE NON-IONIC TRANSPORTERS	Permeability-glycoprotein (P-gp) and multidrug resistant associated protein (MRP).
RECEPTORS	Vitamin B12, vitamin D3, epidermal growth factor and sugar (GLUT1, GLUT3, GLUT5, GLUT2, SGLT1) transports.
DIGESTIVE ENZYMES	Typical membranous peptidases and disaccharidases of the small intestine.

Besides the correlation that exists between *in vivo* drug absorption and Caco-2 permeability [82, 94], the permeability of hydrophilic compounds generally transported by paracellular pathway is low in Caco-2 monolayer because of the relatively tighter junctions compared with human and animal small intestine, which is explained by their colonic origin [95]. Also, the absence of mucus layer will enhance the transport of small molecules due to their molecular characteristics [95]. Furthermore, Caco-2 model cannot differ between cellular transport or intestinal metabolism, since main barriers opposing drug absorption are not completely represented by this model [96].

Despite the widespread use and acceptability of the Caco-2 cell model for permeability assessment, other alternatives have been purposed to overcome Caco-2 limitations (Table 1. 3).

Table 1. 3. Alternatives to *in vitro* drug permeability studies.

CELL LINE	CHARACTERISTICS	REFERENCE
MDCK (Madin Darby Canine Kidney)	<ul style="list-style-type: none"> - Superior model in terms of passive model compared to the Caco-2 model (lower TEER values); - Polarized cells with low intrinsic expression of ABC transporters; - A non-human (dog) and non-intestinal (renal) cell line; - Ideal for transfections; 	[97, 98]
LLC-PK1 (Pig kidney epithelial cells)	<ul style="list-style-type: none"> - Polarized cells with low intrinsic transporter expression; - Ideal for transfections; 	[3]
TC7 (Caco-2 subclone)	<ul style="list-style-type: none"> - Higher levels of CYP3A4 e 3A5; - Useful to evaluate metabolic effects during transport; - Greater homogeneity ensuring more consistent results with lower variability 	[99-101]
2/4/A1 (Rat fetal intestinal cells)	<ul style="list-style-type: none"> - Recommended for passive paracellular transport (leakier monolayers with paracellular pore radius of $9.0 \pm 0.2 \text{ \AA}$, similar to the human small intestine); - TEER values reached a plateau of $50 \Omega\text{cm}^2$, near of the estimated small intestinal TEER values. 	[3, 79, 102]
IEC-18 (Rat intestinal cell line)	<ul style="list-style-type: none"> - Provides a size selective barrier for paracellularly transported compounds. 	[3]

1.7.4. Co-culture models

Caco-2 model lacks in some characteristics that influence drug absorption, for instance, mucus production. With the objective of mimicking the heterogeneity of populations in intestinal epithelium, new Caco-2-based models have been purposed with the introduction of new cell lines.

In this sense, Caco-2 monocultures evolved to co-cultures of Caco-2 and HT29-MTX cells in a similar ratio of the *in vivo* proportions. HT29-MTX cells are a modified cell line treated with an anticancer drug methotrexate (MTX). This cell line came from HT29 cells that exhibit variable differentiation depending on the culture medium changes or differentiation inducers. For instance, when HT29 cells are treated with MTX at 10^{-7} M , they form a mixed population of absorptive and goblet cells. Increased concentrations (10^{-6} to 10^{-5} M) of MTX lead to a population exclusively composed by goblet cells secreting gastric mucins. At higher concentration, is observed a phenotypic change from the mucus to the absorptive phenotype [88].

HT29 cells do not express P-gp and are morphologically distinguished from Caco-2 by the presence of sparse and shorter microvilli, whereas the dimensions of both cells are similar [77]. Caco-2/HT29-MTX model reveals paracellular permeability closer to human situation but the active transport of drugs is underestimated by this model [103]. Interesting works are pointed as proof the concept of this model [103-105].

The promising characteristics of M cells in the induction of specific mucosal immune responses have stimulated the *in vitro* replication of these cells [8]. As result, a new cell culture system consisting of Caco-2 cells and human Raji B lymphocytes aim to establish a model expressing M-like cells originating from Caco-2 conversion by B lymphocytes. This conversion is revealed by the loss of microvilli and consequently by the loss of expression of apical enzymes such as alkaline phosphatase [106-108].

Table 1. 4. Summary of the pros, cons and costs of the intestinal models drug screening and permeability.

	MODELS				
	<i>IN SILICO</i>	<i>EX VIVO</i>	<i>IN VITRO</i>	<i>IN SITU</i>	<i>IN VIVO</i>
	“Rule of five”	Ussing Chamber Everted intestinal sac	Caco-2, HT29-MTX cells and Raji B lymphocytes	Perfusion of drug solution through intestinal segments	Rats, monkeys, dogs and pigs
PROS	<ul style="list-style-type: none"> - Easy access; - High- throughput; - Reduce the time-consuming step of synthesis; - High level of automation. 	<ul style="list-style-type: none"> - Retains gut architecture; - Analysis of different regions of intestine; - The quantity of sample is small and the collected samples are analytically clean; - Possibility of bidirectional transport; - Animal or human models. 	<ul style="list-style-type: none"> - Encompass many characteristics of the intestinal epithelium; - Represents many absorption process; - Can be partly automated; - Information of cell retention; - Benefits in ethical considerations; - Performed under standardized and controlled conditions. 	<ul style="list-style-type: none"> - Presence of an intact blood, nerve supply and clearance capabilities mimic the <i>in vivo</i> conditions; - Evaluation of the intestinal absorption influences the hepatic first pass metabolism; - Good correlation with the pharmacokinetics studies. 	<ul style="list-style-type: none"> - Integration of the dynamic components that affect the drug dissolution; - Relatively to rats, similar absorption profile, low cost and easy care handling. - Monkeys have evolutionary proximity with humans; - Pigs have similarity of large intestine.
CONS	<ul style="list-style-type: none"> - Insufficient data base with reliable data; - Low integration of physiological complexity. 	<ul style="list-style-type: none"> - Tissue viability; - Intrinsic difficult in dissect the epithelial tissue; - Low- throughput; - Requires secondary detection; - Large inter-lab variability; - Difficulty on isolate the contribution of one absorption pathway; - Tissue availability; - The drug must cross the whole intestinal wall. 	<ul style="list-style-type: none"> - Requires secondary detection; - Static model; - Lack in physiological factors (mucus, bile salts); - Genetic drift; - Low differentiation of low/medium absorption; - Large inter-lab variability; - Labor-intensive; - Contamination issues; - Influence of all organism is not taken into account; - Long-differentiation period. 	<ul style="list-style-type: none"> - The method relies on the disappearance of compound from the luminal side, but the rate of decrease of concentration does not always represent the rate of absorption; - Requires a huge number of animals and high amounts of test compounds; - No screening control. 	<ul style="list-style-type: none"> - Data extrapolation must be done carefully due to species differences; - Impact of the anatomical and physiological differences; - Difficulty in separate variables involved in the process of drug absorption; - Ethical concerns; - Extensively resource-intensive. - “The only real model for a man is man.”
COSTS	<ul style="list-style-type: none"> - High development costs. 	<ul style="list-style-type: none"> - High costs per result analysis. 	<ul style="list-style-type: none"> - High in terms of tissue facilities and time-consumable. 	<ul style="list-style-type: none"> - High costs. 	<ul style="list-style-type: none"> - High costs per result analysis.
REFERENCES	[80, 82, 87, 92]	[79, 92, 109]	[21, 88, 91, 96, 104, 105, 110-112]	[113-115]	[83-85, 113]

CHAPTER II

AIM & OBJECTIVES

2.1. Aim and objectives

Caco-2 cell model is the gold standard of intestinal *in vitro* models that after 21 days of culture become differentiated and polarized cells. Their phenotype morphologically and functionally resembles the absorptive enterocytes in many of features, such as, the presence of microvilli, expression of tight junctions, enzymes (alkaline phosphatase, sucrose) and transporters (P-glycoprotein) [112, 116]. However, the value of this model is usually mitigated by tighter tight junctions [91], up-regulation of P-gp [78] and absence of mucus [117]. Gradually, new models have been purposed including Caco-2/ HT29-MTX models [104, 105, 118, 119], Caco-2/Raji B models [21, 120, 121] and Caco-2/HT29-MTX/Raji B models [106-108]. Increasingly, epithelial-stromal interactions have been recognized important players in the maintenance of intestinal architecture. In specific, subepithelial myofibroblasts form a tridimensional network with the underlying cells and play supportive roles promoting epithelial regrowth and enhancement of barrier function [50].

The herein work presents a scenario of interaction between epithelial and stromal cells, replicating the mucosal architecture in such a way that the epithelial layer of Caco-2, HT29-MTX and M-like cells lie over a 3D network of CCD18-Co intestinal myofibroblasts embedded in Matrigel®. The primary aim of this work was to develop a novel intestinal model able to predict the drug permeability of any compound, as closest as the *in vivo* situation. Furthermore, this model has potential to dissect the mutual effect of epithelial and stromal cells in the maintenance of an intact barrier throughout the production of ECM components.

To characterize this model, were established 3 main tasks which aimed to characterize the fibroblastic and epithelial layer, visualize the tridimensional architecture of the model and assess their functional character. Concerning the 21 days of culture and the 3D architecture, CCD18-Co cell density was optimized. Regarding the epithelial layer, the proportion of Caco-2 and HT29-MTX cells was of 9:1 as this ratio replicates the *in vivo* percentage of goblet cells [122] and was previously optimized by our group [107]. The evolution of the barrier model was followed by measuring the transepithelial electric resistance (TEER), evaluating the expression of tight junction proteins between the different epithelial cells, the production of mucus by HT29-MTX cells and the presence of M-like cells. At last, the functional character of the 3D model was assessed by determining the absorption profile of a protein drug, insulin.

CHAPTER III

MATERIALS & METHODS

3.1. Materials

Dulbecco's Modified Eagle's Medium (DMEM), Non-essential aminoacids (NEAA), 0.05% Trypsin-EDTA, 0.4% Trypan Blue, Hanks Balanced Salt Solution (HBSS), Phosphate Buffer Solution (PBS), DMEM without phenol red were purchased from Gibco®, while Fetal Bovine Serum (FBS) and Penicillin/Streptomycin were purchased from BioWest.

Resazurin, Human Insulin, Alcian Blue 8GX, DAPI, Fluoroshield™ mounting medium, Poly-D-Lysine (PDL), 3-aminopropyltriethoxysilane (APES), hexamethyldisilazane (HMDS), Phosphate Substrate (p-NPP) and Propidium Iodide (PI) were purchased from Sigma. 6-well Transwell® permeable support (PET membrane and 3µm pore size) and Matrigel® basement membrane matrix were purchased from BD Biosciences. ClearRite 3, Neg-50™ frozen section medium, paraffin and alcoholic eosin were purchased from Thermo Scientific. Sodium-cacodylate was purchased from Fluka®. Calcein-AM was purchased from Invitrogen. DC Protein Assay Kit was purchased from Bio-Rad. Gill's Haematoxylin II was purchased from Surgipath. Quick-D mounting medium was purchased from Klinipath. Epon resin and propylene oxide were purchased from Electron Microscopy Sciences.

Anti-human vimentin antibody was purchased from Santa Cruz Biotechnology. Donkey anti-mouse AlexaFluor 488®, goat anti-mouse AlexaFluor 594®, goat anti-rabbit AlexaFluor 594®, goat anti-rabbit AlexaFluor 488® secondary antibodies and Quant-iT PicoGreen dsDNA kit were purchased from Molecular Probes®. Rabbit anti-fibronectin polyclonal antibody was purchased from Sigma. Mouse anti-claudin-5 monoclonal antibody was purchased from Life Technologies. Rabbit anti-human E-cadherin (24A10) was purchased from Cell Signaling.

3.2. Establishment of *in vitro* models

3.2.1. Cell lines and cell culture conditions

Caco-2 cell line and HT29-MTX was from American Type Culture Collection (ATCC, USA). Raji B cell line was kindly provided by Dr. Alexandre Carmo (Cellular and Molecular Biology Institute - IBMC, Porto, Portugal). CCD18-Co was kindly afforded by Dr. S rgia Velho (Institute of Molecular Pathology and Immunology of the University of Porto, IPATIMUP, Porto, Portugal).

CCD18-Co (passage 8-13), Caco-2 (passage 22-40) and HT29-MTX (passage 25-40) cells grown separately in tissue culture flasks (Orange Scientific) in DMEM basal medium supplemented with 10% (v/v) of inactivated FBS, 1% (v/v) of NEAA and 1% of antibiotic/antimitotic mixture (final concentration of 100 U/mL Penicillin and 100 U/mL Streptomycin). Cells were routinely observed and, when confluence reached to 80%, subcultured with 0.05% Trypsin-EDTA and seeded at a density of 6×10^5 cells per 75 cm² flask.

The medium was replaced every other day. Raji B cells (passage 30-42) were maintained as aforementioned and seeded at a density of 1×10^6 cells per 75 cm^2 flask.

Cells were maintained in an incubator (CellCulture[®] CO₂ Incubator, ESCO) at 37°C and 5% CO₂ in water saturated atmosphere. All the cell culture related procedures were done in a Labculture[®] class II, type A2 biological safety cabinet, ESCO.

3.2.2. Cell thawing

All cells were frozen at liquid nitrogen. It is essential to thaw the cell correctly in order to maintain the viability of the culture and allow them to recover quickly. Furthermore, some cryoprotectants, such as DMSO, are toxic above 4°C so cultures should be thawed quickly and diluted in culture medium to minimize the toxic effects. Despite this, DMSO protects the cells from rupture by the formation of ice crystals.

The basic principle of a successful resuscitation is a quick thaw, by incubation in a 37°C water bath for 3-4 minutes. All the content in the cryovial (CryoVial[®], Falcon) was transferred to a new container with pre-warmed culture medium and centrifuged at 18 G for 5 minutes at 4°C to remove at DMSO. The supernatant was discarded and the pellet was resuspended in pre-warmed complete DMEM. The content was transferred to a tissue culture flask (Orange Scientific) and cell viability was evaluated after 24 hours.

3.2.3. Optimization of CCD18-Co cells density

To determine the optimal density of CCD18-Co intestinal myofibroblasts, different cellular densities of 1×10^4 cells/cm², 5×10^4 cells/cm², 10×10^4 cells/cm² and 20×10^4 cells/cm² were evaluated regarding their viability, metabolic activity, proliferation and morphological features.

3.2.3.1. Cell viability

At different time points, CCD18-Co cells were washed twice with pre-warmed DMEM without phenol red to remove/dilute estereases and incubated with Calcein-AM diluted in DMEM without phenol red (1:250) for 40 minutes at 37°C. After washed twice with DMEM without phenol red, samples were incubated with PI diluted in DMEM without phenol red (1:250) for 10 minutes at 37°C. Samples were finally washed twice with DMEM without phenol red and immediately examined under an Axiovert 200M inverted fluorescent microscope from Zeiss. Live cells are distinguished by the presence of ubiquitous intracellular esterase activity that is able to digest non-fluorescent calcein-AM into fluorescent calcein emitting green fluorescence [111]. In contrast, dead cells with damaged membranes allow the entrance of propidium iodide (PI) that emits a bright red fluorescence upon binding to the acid nucleic.

3.2.3.2. Cell metabolic activity

Metabolic assays were conducted using the resazurin fluorescent assay. Representative of 3D condition, CCD18-Co cells were embedded Matrigel® matrix, while other set was prepared by seeding CCD18-Co, case of study representative of 2D standard conditions.

Resazurin pre-warmed to 37°C was added to the plates in a 10% concentration of the initial volume in the plate and cells in culture for 7 days were incubated at 37°C. After incubation, a sample was collected and immediately quantified by measuring the relative fluorescence units (RFU) using a microplate reader Synergy™ Mx HM550 (BioTek® Instruments, USA) set at 530/590 nm (excitation/emission wavelength, respectively). The results were normalized by subtracting the negative control (without cells). The assay was repeated for 14 and 21-days cultures.

In this assay, the cellular metabolic activity is detected through the reduction of a blue non-fluorescent resazurin reagent into a highly fluorescent resorufin by the dehydrogenase enzyme in metabolically active cells. This conversion is proportional to the number of viable cells in the sample.

3.2.3.3. Total dsDNA quantification

After recover the cells by trypsinization, the cell suspension was centrifuged at 18 G for 5 minutes, washed with PBS, centrifuged again and the pellet was stored at -20°C until quantification. Cells were lysed with 1% (v/v) Triton X-100 for 1 hour under agitation at 4°C. Since 1% (v/v) Triton X-100 interferes with the measurement, samples were diluted to a final concentration of 0.4% (v/v) Triton X-100 with PBS and the double-stranded DNA (dsDNA) was quantified using the Quant-iT PicoGreen dsDNA kit, according to manufacturer's instructions. Briefly, 10 µL of each sample was transferred to a black 96-well plate with clear bottom (Falcon) and diluted with 90 µL of TE (1x) buffer (200mM Tris-HCl, 20 mM EDTA, pH 7.5). To each well, was added 100 µL of PicoGreen dsDNA reagent and incubated for 5 minutes at RT, protected from light. Samples were quantified using a microplate reader Synergy™ Mx HM550 (BioTek® Instruments, USA) set at 480/530 nm (excitation/emission wavelength, respectively). The dsDNA concentration (ng/mL) was determined using a standard curve of lambda DNA in a range of 0.0250-2.50 ng/mL.

3.2.3.4. Cell morphology

CCD18-Co cells morphology was assessed by labeling the intermediate filaments of fibroblasts with vimentin antibody.

Briefly, CCD18-Co cells, previously seeded on coverslips coated with PDL, were rinsed twice with pre-warmed PBS⁺ (1.37 M NaCl, 26.8 mM KCl, 14.7 mM KH₂PO₄, 78.1 mM Na₂HPO₄, 1mM Ca₂Cl, 1mM Mg₂Cl, pH 7.4), regular PBS supplemented with Ca²⁺ and Mg²⁺ ions to

maintain the integrity of cellular junctions. Cells were fixed with 2% (w/v) paraformaldehyde (Merck, VWR) in PBS⁺ for 15 minutes at room temperature (RT), following permeabilization with freshly prepared solution of 0.2% (v/v) Triton X-100 (Sigma®, Sigma Aldrich) in PBS⁺ for 7 minutes at RT. In order to block non-specific binding, samples were incubated with a freshly prepared solution of PBST (PBS⁺ containing 0.05% (v/v) Tween-20 (AppliChem, VWR)) containing 10% (v/v) FBS for 30 minutes in a humidified chamber. Then, without wash, blocking solution was removed and samples were separately incubated during 2 hours at RT with a mouse anti-human vimentin primary antibody diluted (1:100) and a donkey anti-mouse AlexaFluor488® secondary antibody (1:1000) during 1 hour at RT, both diluted in PBST containing 5% (v/v) FBS. Cell nucleus was counterstained with DAPI diluted 1:1000 in PBS during 5 minutes. With exception of the blocking step, were performed 5 minutes washes with PBS⁺ between fixation, permeabilization and incubation steps. The coverslip was placed face down to the glass slide with a drop of mounting medium of Fluoroshield™ and visualized using an Axiovert 200M inverted fluorescent microscope from Zeiss.

The acquired images were taken using Axiovision 4.8.2 SP3 from Zeiss and treated using ImageJ program. The staining was repeated for 14 and 21 days old cultured cells to understand the effect of the initial cell density on fibroblasts morphology.

3.2.4. Production of ECM components

CCD18-Co cells entrapped in Matrigel were evaluated regarding expression of ECM components, namely fibronectin. Moreover, the 3D arrangement of CCD18-Co cells entrapped in the Matrigel matrix was also assessed. The procedure was performed as previously reported for paraffin sections with exception of the incubation step. Briefly, cells were incubated during 2h at RT with both mouse anti-human vimentin primary antibody (1:100) and rabbit anti-human fibronectin primary antibody (1:400). After 5 minutes washes, a goat anti-mouse AlexaFluor594 secondary antibody (1:1000) and a goat anti-rabbit AlexaFluor488 secondary antibody (1:1000) were incubated during 1 hour at RT and protected from light. Cell nucleus was counterstained with DAPI diluted 1:1000 in PBS during 5 minutes.

Paraffin-embedded sections of the 3D model (CCD18-Co+Matrigel/Caco-2/HT29-MTX/Raji B) were stained to evaluate the supportive role of fibroblasts embedded in self-produced extracellular matrix. First, sections were deparaffinise with ClearRite for 10 minutes and hydrated through graded series of ethanol finishing with PBS (1x), during 5 minutes in each solution. Sections were fixed with 4% (w/v) paraformaldehyde (Merck, VWR) in PBS⁺ for 20 minutes at RT, following a permeabilization with freshly prepared solution of 0.2% (v/v) Triton X-100 (Sigma®, Sigma Aldrich) in PBS⁺ for 9 minutes at RT. The blocking was performed with 1% BSA in PBS⁺ during 30 minutes at RT. Then, without wash, blocking solution was removed and sections were incubated during 2 h at RT with a rabbit anti-human fibronectin primary antibody (1:400) and a goat anti-rabbit AlexaFluor488 secondary antibody

(1:1000) during 1 hour at RT, both diluted in 1% BSA in PBS⁺. Cell nucleus was counterstained with DAPI diluted 1:1000 in PBS during 5 minutes. With exception of the blocking step, were performed 5 minutes washes with PBS⁺ between fixation, permeabilization and incubation periods.

Samples were mounted with Fluoroshield™, visualized using an Axiovert 200M inverted fluorescent microscope from Zeiss and treated using ImageJ program.

3.2.5. Monolayer *in vitro* models

As manner of proof, single and co-culture monolayers were performed comprising Caco-2, HT29-MTX, CCD18-Co and Raji B cells. By single monolayers are intended Caco-2 and CCD18-Co monolayer's while co-culture monolayers involve CCD18-Co/Caco-2 cells, Caco-2/HT29-MTX cells, Caco-2/Raji B cells and Caco-2/HT29-MTX/Raji B cells.

Stromal - epithelial co-culture of CCD18-Co/Caco-2 cells (**Fig. 3. 1 A**) was prepared by seeding CCD18-Co at a density of 1×10^4 cells/cm² onto 3 μ m Transwell™ inserts. In the following day, Caco-2 cells were seeded onto CCD18-Co cells at a density of 1×10^5 cells/cm², as described before [106, 107].

Epithelial co-culture of Caco-2/HT29-MTX cells (**Fig. 3. 1 B**) was prepared by seeding Caco-2 and HT29-MTX cells to a final density of 1×10^5 cells/cm² (90:10 proportion), previously optimized [107]. These cultures were maintained for further 21 days.

The epithelial - lymphocyte co-culture of Caco-2/Raji B cells (**Fig. 3. 1 C**) was prepared by seeding Caco-2 at a final density of 1×10^5 cells/cm² in the apical side of the Transwell™ insert, while 1×10^6 cells/6-well Raji B lymphocytes were added to the basolateral side after 14 days in culture. Until the end of 21 days, medium was changed every other day only at the apical compartment.

The triple model of Caco-2/HT29-MTX/Raji B cells (**Fig. 3. 1 D**) was performed as described before [106, 107] based on the Caco-2/HT29-MTX co-culture with the addition of Raji B cells in the basolateral side.

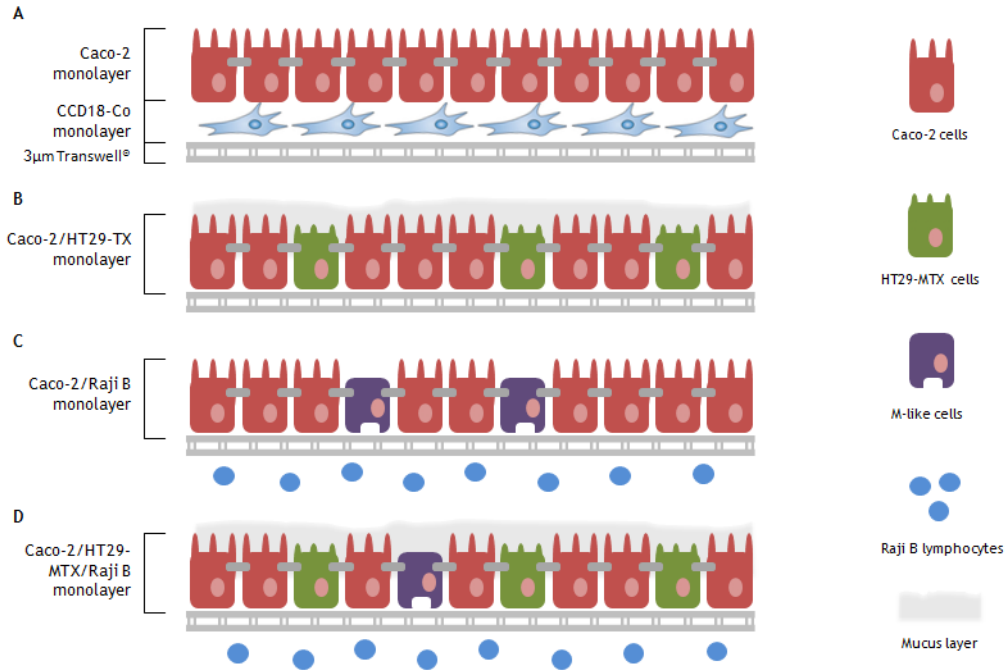


Figure 3. 1. Schematic configuration of monolayer *in vitro* models. (A) Stromal - epithelial monolayer comprising CCD18-Co and Caco-2 cells. (B) Epithelial monolayer comprising Caco-2 and HT29-MTX cells. (C) Epithelial - lymphocyte monolayer representative of the formation of M-like cells. (D) Triple model comprising Caco-2, HT29-MTX and M-like cells.

3.2.6. Tridimensional *in vitro* models

To explore the impact of CCD18-Co cells' assembly in the barrier formation were tested distinct configurations. The first configuration consists on Caco-2 cells beneath CCD18-Co cells embedded in Matrigel (CCD18-Co cells in 3D assembly) (**Fig. 3. 2 A**) and the second configuration consists on CCD18-Co and Caco-2 cells separated by a gel layer of Matrigel between both cell types (CCD18-Co cells in 2D assembly) (**Fig. 3. 2 B**).

For the first configuration, 1×10^4 cells/cm² of CCD18-Co cells were entrapped in a thick layer (400µL) of Matrigel (1:3) diluted in ice-cold DMEM without FBS. The cell suspension was spread evenly by the Transwell® insert from the centre to the periphery, avoiding bubbles. The Transwell® insert was transferred to the incubator at 37°C and allowed to polymerize for 30 minutes. After that, DMEM was gently added to both sides of the Transwell® insert. After 24h, 1×10^5 cells/cm² of Caco-2 cells were added over the layer of CCD18-Co embedded in Matrigel®.

For the second configuration, CCD18-Co cells were seeded to a final density of 1×10^4 cells/cm². In the following day, DMEM as removed and a thin layer (200µL) of Matrigel® (1:3) was added onto the CCD18-Co monolayer. After 30 minutes of polymerization, Caco-2 cells were carefully added over the Matrigel® layer. The cell density of CCD18-Co cells was based on preliminary studies and both culture models were maintained for further 21 days.

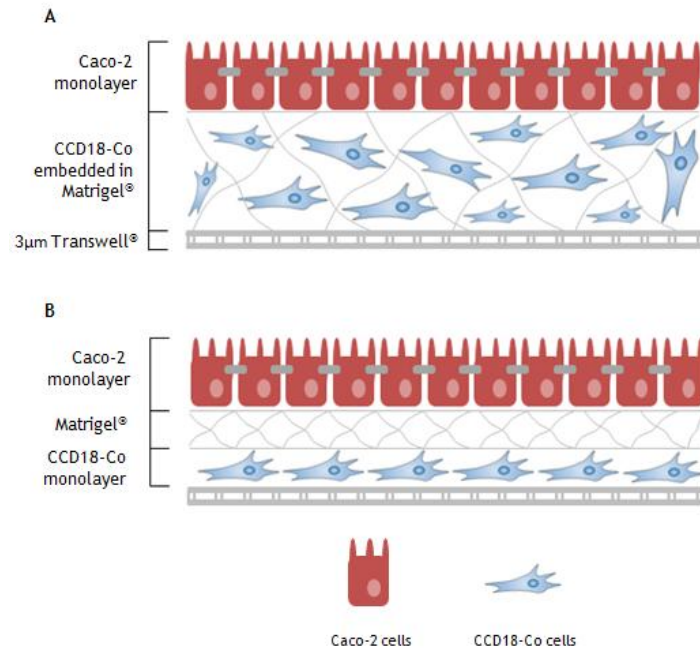


Figure 3. 2. Schematic configuration of the double 3D *in vitro* models. (A) CCD18-Co cells embedded in a Matrigel® matrix, representing a 3D assembly. (B) CCD18-Co cells seeded on the bottom of the Transwell, representing a 2D assembly.

To advance to further studies was adopted the first configuration based on the preliminary studies obtained. Moreover, this arrangement closely mimics the intestinal mucosa. The final model, which adds HT29-MTX and M-like cells converted by Raji B cells, is represented in Fig. 3. 3.

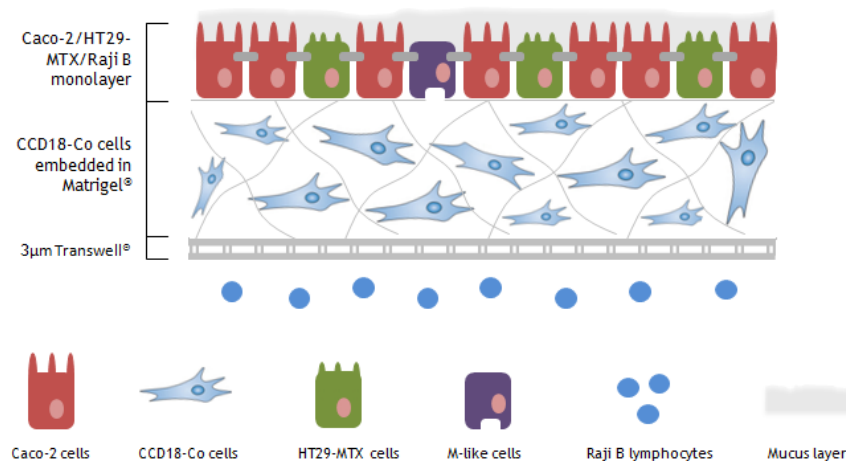


Figure 3. 3. Schematic configuration of the 3D quadruple model.

3.2.7. Transepithelial electrical resistance (TEER)

To monitor the membrane integrity, the transepithelial electrical resistance (TEER) of each model was measured using a voltohmmeter Millicell® ERS-2 (Millipore, USA) every time medium was replaced. The values were normalized by subtracting the resistance value of the empty Transwell® insert and expressed in ohm per area of the insert ($\Omega \cdot \text{cm}^2$) through the following equation.

3.2.8. Cell proliferation

The effect of the Matrigel® on cell proliferation of stromal and epithelial cells was explored by comparing the behaviour of CCD18-Co cells at a density of 1×10^4 cells/cm² and Caco-2 at a density of 1×10^5 cells/cm² in plastic and Matrigel. CCD18-Co cells were also seeded on plastic in direct contact with Caco-2 cells.

After 7 days, cells were washed twice with PBS (1.37 M NaCl, 26.8 mM KCl, 14.7 mM KH₂PO₄, 78.1 mM Na₂HPO₄, pH 7.4) and detached from substrates with 0.05% trypsin-EDTA during 10 minutes at 37°C. Trypsin-EDTA mixture was inactivated with DMEM and the plates were observed to the microscope to assure that all cells were detached; otherwise trypsinization was repeated. Each sample was mixed with 0.4% (v/v) Trypan Blue to determine the cell number using a haemocytometer (Marienfeld-Superior, Germany).

3.2.9. Paraffin sections and histological staining

Membranes were embedded in paraffin to observe the layers that constitute the 3D model. Briefly, membranes were washed twice with PBS before fixed with 4% paraformaldehyde for 1 hour and post-fixed with 1% glutaraldehyde for 15 minutes to avoid depolymerization of Matrigel. Membranes were washed twice with PBS and cut with a scalpel to be processed automatically in a Spin Tissue Processor STP120 (Thermo Scientific, Germany) that consisted in graded dehydrations in ethanol for 10 minutes each, followed by a diaphanization using ClearRite 3 for 10 minutes twice. Membranes processing was finished by placing membranes in liquid paraffin for 2 hours. Membranes were orientated according the plane of cut and sectioned with a thickness between 5-10 μm using a Leica RM2255 microtome (Leica Biosystems, Germany). The glass slides with the sections were allowed to dry overnight at 37°C before haematoxylin and eosin staining.

Briefly, for the haematoxylin and eosin staining, sections were dewaxed in ClearRite 3 for 10 minutes. After that, sections were hydrated through graded alcohols to water for 2 minutes each, following the staining with Gill's haematoxylin II for 5 minutes. Sections were abundantly washed in tap water for 2 minutes, following dehydration through graded alcohols for 2 minutes each. Sections were stained with alcoholic eosin for 3 minutes and sections

were dipped in absolute alcohol 3 quick times. At last, sections were dipped in xylene three times and mount in Quick-D permanent mounting medium.

3.2.10. Expression of junctions proteins

Immunofluorescence staining was used to determine the expression of the TJ proteins claudin-5 and occludin and adherent junction protein E-cadherin.

Regarding claudin-5 labeling, cells were washed twice with pre-warmed PBST for 1 minute each. After that, cells were fixed with 2% (w/v) PFA for 15 minutes at RT following the permeabilization step with 0.2% of Triton during 5 minutes. After washes, the non-specific binding was blocked with PBST containing 10% FBS for 30 minutes. Then, without wash, the mouse anti-claudin-5 conjugated with AlexaFluor 488 antibody was incubated during 1 hour in a 1:100 dilution in PBS. Samples were washed with PBS⁺ and incubated with DAPI (1:1000) for 5 minutes to label the cell nuclei.

Regarding the occludin labeling, cells were washed with cold PBS⁺ and fixed for 10 minutes on ice with methanol. After this, cells were washed 3 times for 5 minutes with PBS⁺ and non-specific binding was blocked incubating with 5% (v/v) BSA at RT. Then without wash, mouse anti-occludin conjugated AlexaFluor 594 antibody was incubated for 1 hour at RT and protected from light, in a dilution of 1:100 in PBS. Samples were washed with PBS⁺ and incubated with DAPI (1:1000) for 5 minutes to label the cell nuclei.

Concerning the E-cadherin labeling, samples were washed with PBS⁺ that allows the maintenance of junctions, following fixation with 4% (w/v) PFA for 20 minutes at RT. Samples were incubated with 50 mM NH₄Cl (Sigma®) for 10 minutes that blocks aldehyde groups allowing a better antibody-antigen bonding. After permeabilization with 0.2% Triton for 5 minutes, the non-specific binding was blocked with 5% BSA for 30 minutes. The blocking solution was removed and samples were incubated with the rabbit anti-human E-cadherin primary antibody diluted to 1:100 in 5% BSA for 2 hours. It follows the incubation with the goat anti-rabbit AlexaFluor488 secondary antibody diluted to 1:500 in 5% BSA. The staining was finalized with nuclei labeling with DAPI (1:1000). Samples were mounted with Fluoroshield™ and visualized using an Axiovert 200M inverted fluorescent microscope from Zeiss.

3.2.11. Alkaline phosphatase assay

Alkaline phosphatase activity was measured *in situ* live cells according to Ferruza *et al.* using p-Nitrophenyl (p-NPP) as substrate [123]. Alkaline environment catalyses the hydrolysis of p-NPP to a inorganic phosphate and p-Nitrophenol (p-NP), a yellow soluble end-product. The intensity of the colored product is proportional to the enzyme activity measured at pH 9.3 at 405 nm.

Briefly, after removing the cell medium and gently wash the monolayer with PBS⁺ (1mM CaCl₂ and 1mM MgCl₂), reaction buffer consisting of one volume of collection buffer (10 mM Tris-HCl, 150 mM NaCl pH 8.0) and three volumes of pNPP solution (100 mM Tris-HCl, 100 mM NaCl, 5 mM MgCl₂, pH 9.5) was added to the well, kept at 37°C in water bath. At different time points, 100 µL was collected and transferred to a 96-well plate containing 50 µL of 0.5M of NaOH to stop the reaction. The standard curve was prepared with 50 µL of each p-NP concentration ranging from 0 to 600 µM, added to 150 µL of p-NPP solution and 50 µL of 0.5M NaOH. Absorbance was determined using a SynergyTM Mx HM550 microplate reader (BioTek[®] Instruments, USA) and normalized by blank subtraction. Concentration was determined with reference to the standard curve and corrected for the reaction volume before plotting was function of the time. The angular coefficient of the linear slope of this plot represents the enzymatic activity expressed as mU= nmol of p-NP/min at 37°C and normalized to mg of cellular protein measured Bio-Rad DC Protein Assay, a Lowry-based assay [124].

3.2.11.1. Protein quantification

After the alkaline phosphatase assay, samples were lysed for further protein quantification using Bio-Rad DC Protein Assay, which is equivalent to the well documented Lowry assay. This assay is based in two steps that lead to the color development. First, the reaction between the protein and the copper in an alkaline medium (Reagent A) and subsequent reduction of Folin Reagent (Reagent B) by the copper treated protein [124].

The assay was performed according to the manufacturer's instructions. Briefly, cells were incubated with the lysis buffer (250mM Tris and 0.1% Triton) for 20 minutes at RT. The cell suspension was collected and was added 1 volume of dilution buffer (250mM Tris) to reduce the amount of detergent. Samples were vortex and frozen until further quantification. For quantification purposes, samples were centrifuged at 18 G for 5 minutes to pellet the insoluble material and 5µL of each standard and samples was transferred to a 96-well plate, at which was added 25 µL of Reagent A' (20 µL of Reagent S to each mL of Reagent A) and 200 µL of Reagent B. The plate was mixed for 5 seconds and after 15 minutes, absorbance was read at 750 nm. Concentration was determined with reference to a standard curve of BSA between 1.5 mg/mL and 0.125 mg/mL.

3.2.12. Scanning electron microscopy

After 21 days in culture, Transwell membranes were washed twice in PBS⁺, following primary fixation with 3% (v/v) Glutaraldehyde (AGAR Scientific, AGAR Scientific) for 45 minutes. After rinsing with 0.1M sodium-cacodylate buffer, membranes were dehydrated through a graded series of ethanol, subsequently dried with 75% (v/v) hexamethyldisilazane (HMDS) in ethanol for 10 minutes. Without remove HMDS, Transwell inserts were allowed to air-dry for 30 minutes. The membrane of the Transwell inserts was

removed from the base of the insert using a scalpel and mounted on double-side stick tape SEM stubs. The samples were stored in a desiccator until coated. Sputter coat was with gold-palladium at 15 mA for 120 seconds under argon atmosphere and images were acquired using a scanning electron microscope (Quanta 400FEG ESEM/EDAX Genesis X4M).

3.2.13. Transmission electron microscopy

After 21 days in culture, Transwell membranes were washed twice in PBS⁺, following primary fixation with 2.5% (v/v) Glutaraldehyde (AGAR Scientific, AGAR Scientific) in 0.1M sodium cacodylate during 3 days at 4°C. After rinsing with 0.1M sodium-cacodylate buffer, membranes were post-fixed in 2% (v/v) osmium in 0.1M sodium cacodylate buffer overnight. Membranes were washed with 0.1M sodium-cacodylate buffer, following dehydration through a graded series of ethanol and propylene oxide for 10 minutes each. The membrane was infiltrated in embedding resin as follows: propylene oxide and epon (2:1 proportion), 1:1, 1:2 and 0:1 during 1 hour each. After that, resin was allowed to harden during 48 hours in an incubator set at 65°C. The block was sectioned with 60 nm of thickness using a Leica Reichert SuperNova ultramicrotome (Leica Microsystems, Germany) and samples are collected to an aluminum disk. The disks were contrasted using uranyl acetate and lead citrate during 5 minutes in each solution. Images were acquired using a transmission electron microscope Jeol JEM 1400.

3.2.14. Drug transport assay

In order to determine the permeability of model drugs through the developed cell-based models, permeability studies were performed correlating the obtained values with the *in vivo* data. To start, culture medium was removed from both chambers and the cells membranes were washed with pre-warmed HBSS, further replaced by new HBSS and allowed to equilibrate for 15 minutes at 37°C. After this period it is expected that cell cultures reach a constant TEER value. Permeability studies were run at 37°C during 4 hours with 1.5 mL of 100 µg/mL insulin prepared in HBSS in the apical side and 2.5 mL of HBSS in the basolateral side.

Cell monolayer integrity during permeability studies was controlled by TEER measurement before collect the samples in different time points. The samples were further quantified by reverse HPLC-UV on a Merck-Hitachi LaChrom HPLC instrument (Merck). The HPLC system was equipped with a XTerra RP-18 column, 5 µm particle size, 4.6 mm internal diameter × 250 mm length (Waters) and a LiChrospher 100 TP-18, 5 µm particle size guard column (Merck) [125]. All experiments were run in triplicate at room temperature and the total area of the peak was used to quantify insulin. The results were expressed in percentage of release and apparent permeability (P_{app}). P_{app} was calculated using the following equation:

$$P_{app} = \frac{\Delta Q}{A \times C_0 \times \Delta t} ,$$

, where C_0 is the initial concentration in the apical compartment ($\mu\text{g/mL}$), A is the surface area of the insert (cm^2), Δt is the time in which the experiment occurred (seconds) and ΔQ is the amount of compound detected in the basolateral side (μg) [107].

Complementary studies to integrity were performed at the end of permeability studies to evaluate the final integrity state of the models. For this, apical-basolateral flux was assessed using 4 kDa Fluorescein Isothiocyanate - Dextran molecule (FITC-Dextran). After transport experiments with insulin, it was discarded and Transwell® inserts were washed twice with HBSS. After washes, 1.5 mL of 200 $\mu\text{g/mL}$ of FITC-Dextran was added to the apical compartment, equilibrated with 2.5 mL of HBSS in the basolateral side. The plate was incubated at 37°C, 5% CO_2 for 1 hour. After 1 hour, a sample was collected from the basolateral side and transferred to a black plate 96-well plate (Falcon). FITC-Dextran quantification was determined by fluorescence using a Synergy™ Mx HM550 microplate reader (BioTek® Instruments, USA) set at 485/530 nm (excitation/emission wavelength, respectively). Microplate readings were obtained through Gen5 Data Analysis software (BioTek® Instruments). Fluorescence values were normalized subtracting the blank and concentration was calculated with reference to the standard curve. The amount of transported 4 kDa FITC-Dextran was expressed by percentage of rejection.

$$\% \text{ Rejection} = \frac{1 - RFU_{basolateral}}{RFU_{apical}} \times 100$$

3.2.15. Statistical analysis

All the results are presented as the mean \pm SD. Statistical analysis was performed using Graphpad Prism® software (Prism 6 for Windows, CA, USA).

One way ANOVA, together with Tukey's post hoc test, was used to compare different groups of data. $P < 0.05$ was considered statistically significant.

CHAPTER IV

RESULTS & DISCUSSION

The growing interest in predicting the properties of incoming drugs in the market has increasingly valuing the *in vitro* tools as they may encompasses many characteristics of the tissue in question and also benefit from standardized conditions and lack of ethical considerations. Nonetheless, the existing *in vitro* models are restricted to cell monolayers that do not replicate the entire architecture and mechanisms occurring in a living tissue [126]. Essential cellular functions are missed by ‘petri-dish’-based cell cultures, limiting the potential to predict the cellular responses of real organisms. Tridimensional cultures are believed to incite a strong impact on drug screening, bridging the gap between cells cultures and animal models, with the potential to additionally decrease the use of animals in research [127].

The present study extends a traditional standard intestinal model to the deeper layers of small intestine, in particular the intestinal mucosa that includes a broad variety of stromal cells. The intimate crosstalk between epithelial and stromal cells was explored using a novel *in vitro* cell based model as a mean of mimic the intestinal mucosa’s architecture and functioning and predicts the permeability of a model drug.

4.1. Analysis of CCD18-Co cells behaviour in 2D and 3D

There is a striking similarity between the morphology and behaviour of cells *in vivo* and in 3D culture conditions [127]. The cell behaviour in 2D and 3D matrices has revealed considerable differences in cell morphology [128], migration [129], adhesion [130] and signalling [131], comparing to *in vivo* conditions. Together, these findings empower the importance of matrix dimensionality, i.e. 2D vs. 3D.

The evaluation of cell behaviour in 3D microenvironments is a multifactorial task that takes into account matrix extrinsic factors (soluble and direct cell-cell contact signaling) and matrix related factors (matrix composition, mechanical properties, mesh size and degradation rate) [132]. A recent study highlights the relevance of optimizing initial cell densities when establishing 3D cultures [133].

To determine the optimal cell number of CCD18-Co cells intestinal myofibroblasts several assays were performed, namely viability, metabolic activity and cell morphology over 21 days of culture, period necessary to fully differentiate Caco-2 cells [93]. In this sense, a range of densities of CCD18-Co cells (between 1×10^4 cells/cm² (D1) and 20×10^4 cells/cm² (D4)) was seeded on 24-well plates to assess the most suitable density to be used in further experiments.

4.1.1. Effect of cell number on viability of CCD18-Co cells in 2D

Cell viability was evaluated at days 7, 14 and 21 by the Live/Dead assay. As observed in Fig. 4. 1, after 7 days, the cells remained highly viable for all the range of densities, with very few dead cells between them. Despite the high viability at day 14, an alteration of cell

shape at density D3 (10×10^4 cells/cm²) and D4 (20×10^4 cells/cm²), can already be observed since calcein is ubiquitously distributed by the cells. Cells seem to acquire a starry-like shape most likely as a result of space constraints at higher cell densities. It is known that high density cell culture promotes cell-to-cell contact that can stimulate cell cycle arrest causing cells to stop dividing [134], a phenomenon commonly known as contact inhibition [135].

The starry-like shape is more accentuated at day 21, indicating that D3 and D4 do not seem appropriate cell densities using 2D standard culture conditions. Notwithstanding, further morphological assays were carried out to sustain this result.

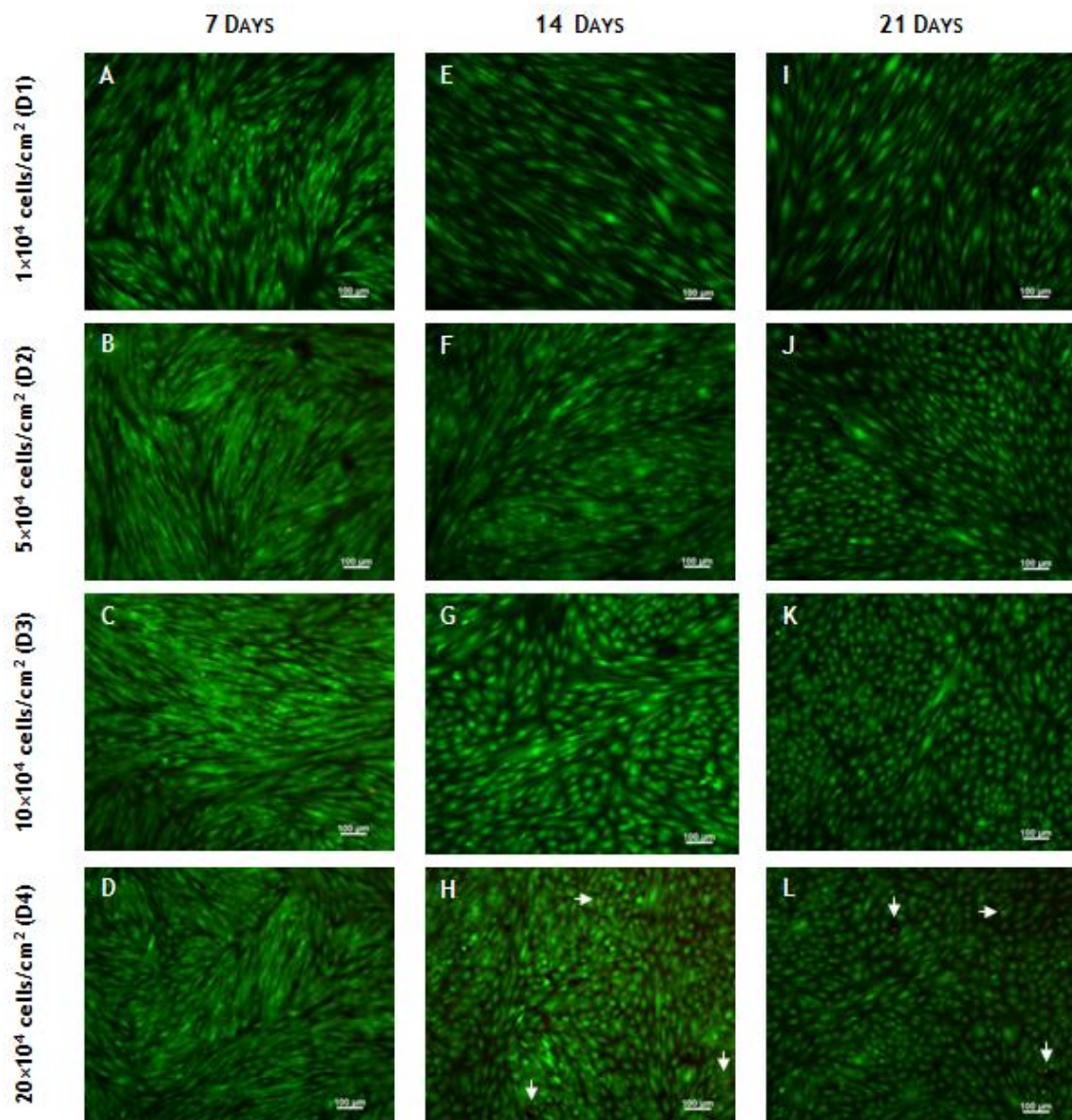


Figure 4. 1. Live/Dead viability assay performed in 2D cultures of CCD18-Co cells. The results showed that at day 7 (A-D) most of the cells are viable labeled with Calcein-AM (green) presenting an elongated shape. After 14 days (E-H), cells at higher densities (D3 and D4) start to acquire a starry-like shape with few dead cells labeled with PI (red). This tendency aggravates at day 21 (I-L) for lower densities (D2). Scale indicates 100 µm and white arrows indicate dead cells.

4.1.2. Effect of cell number on metabolic activity of CCD18-Co cells

The metabolic activity of CCD18-Co cells was evaluated in 3D conditions after entrapment in a matrigel matrix and in standard 2D conditions using the rezasurin assay. The differences in cell behaviour were evaluated during 21 days. Despite the different culture conditions, the time of incubation for both 2D and 3D cultures was 3 hours.

In 2D standard conditions (Fig. 4. 2 A), the metabolic activity presented a stairway profile between D1 and D3 for 7 days. At day 14, an increase in metabolic activity was observed for all the cell densities tested, although a growing approximation of the fluorescent values between D2, D3 and D4 was noticed. At day 21, the metabolic activity decreased for all the densities, probably result of contact inhibition [135]. Concerning the 3D cultures (Fig. 4. 2 B), the metabolic activity also presented a stairway profile for the first 7 days, proportional to the cell densities. At day 14, the increase of metabolic activity at D4 was not significant and after 21 days, 3D cultures seemed to acquire a stagnant profile.

Overall, in the first days of culture, the metabolic activity of 3D cultures was lower when compared with 2D cultures. Under 3D conditions, cells have a limited proliferation as the matrigel matrix constitutes an obstacle to the proliferation and cells are subject to hypoxia. But in fact, after 21 days, the metabolic activity of 3D cultures was higher when compared with 2D cultures. This probably indicates that to cells maintain metabolically viable they had used their proteolysis machinery and migrated through the matrix [136]. For this reason, the choice of matrigel dilution and thickness are important parameters that will condition the cell behaviour as the pore size of the matrigel matrix varies with the protein concentration. At 50% gel concentration, the average pore size is approximately 2 μm , which is much smaller than cellular dimensions [136]. Even though balanced matrix degradation allows cell survival throughout matrix remodeling, further studies are necessary to evaluate the matrix degradation and production of metalloproteinases (MMPs).

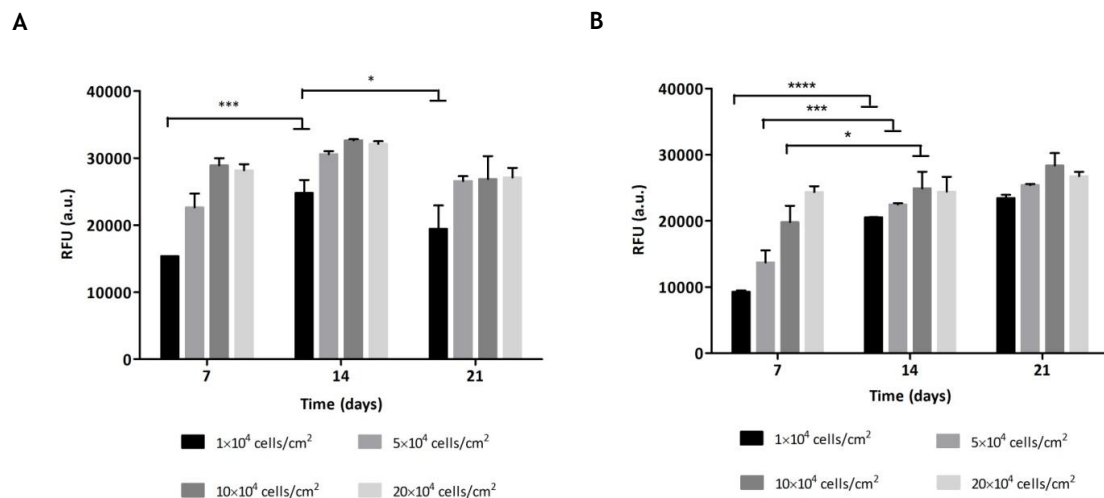


Figure 4. 2. Metabolic activity of CCD18-Co cultured in 2D and 3D along 21 days at different densities. The metabolic activity of 3D cultures was higher compared with 2D cultures after 21 days. Results are the average of replicates and bars represent the standard deviation (n=3). The statistical differences ($p < 0.05$) are denoted by (*).

4.1.3. DNA quantification

As mean to reinforce the results obtained for the metabolic activity in 2D and 3D cultures, it was determined the DNA content after 21 days of culture (Fig. 4. 3). At day 1, it was observed that the DNA content was proportional to the range of cell densities tested. After 21 days, the DNA content in 2D cultures was significantly reduced at D2, D3 and D4 but not at D1. This means that at D1, cells did not enter in a process of contact inhibition. Therefore, cell densities higher than 1×10^5 cells/cm² (D2) are not recommended for cultures maintained for 21 days in 2D standard conditions. Cells start to lack of space, detaching and being removed as the medium is replaced every other day.

The decrease in the DNA content observed in 3D is a result of the reduced proliferation of cells when are entrapped in a matrix. Notwithstanding, the DNA content was proportional to the cell density.

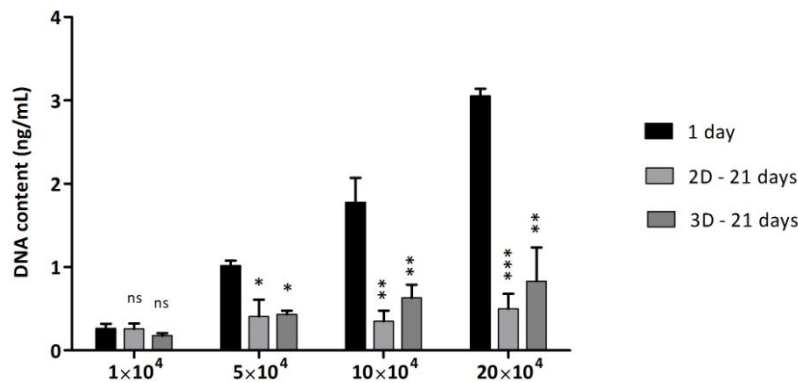


Figure 4. 3. Total dsDNA quantification of CCD18-Co cultured for 21 days in 2D standard conditions at different densities. After 21 days, the decrease in DNA content in both 2D and 3D cultures was result of contact inhibition and matrix constraints, respectively. Results are the average of replicates and bars represent the standard deviation (n=3). The statistical differences ($p < 0.05$) are denoted by (*) regarding the control at day 1.

In summary, when entrapped in a matrigel matrix, CCD18-Co cells revealed a non-proliferative character, result of the matrix constraints.

4.1.4. Effect of cell number in CCD18-Co cells morphology

There are, at least, three major classes of microenvironmental factors that affect the cellular behaviour, i.e., the biochemical composition, the physical parameters of the matrix, particularly the pliability, and the spatial cues [137]. 2D surfaces do not represent the complexity of factors involved in cell communication and growth. However, growth arrest and consequent metabolic activity inhibition are a consistent issue in 3D matrices added to the fact that cells are spatially constrained and need to overcome the environment impediments to populate the surrounding environment [138].

Regarding these issues, the morphology of CCD18-Co cells under 2D culture conditions was evaluated to further on compare with the cell morphology in 3D culture conditions. After

7 days were observed remarkable differences between cells densities, highlighting the elongated cell shape only at the density 1×10^4 cells/cm² (Fig. 4. 4 A). Cell morphology suffered abrupt alterations beyond 1×10^4 cells/cm² cell density, result of the contact inhibition and the lack of space. Between 5×10^4 cells/cm² and 10×10^4 cells/cm², cells adopted a starry-like morphology, still mixed with some elongated cells (Fig. 4. 4 B and C). For the density of 1×10^5 cells/cm², the population of cells with the characteristic fibroblast shape was found on odd occasion. Distinctly, at the highest density (20×10^4 cells/cm²), CCD18-Co cells presented a skinny cell body with some cells overlapping in some regions (Fig. 4. 4 D).

At 14 days, the previous scenario is aggravated but 1×10^4 cells/cm² cell density seems to be the better adequate density presenting a good spatial organization and maintained the original and characteristic morphology of myofibroblasts (Fig. 4. 5 A). At higher densities (D2 and D3), the starry-like population of fibroblasts is more prominent (Fig. 4. 5 B) observing the existence of a syncytium at D3 (Fig. 4. 5 C), a multinucleated cell that can result from multiple cell fusions of mononuclear cells. The existence of a syncytium is reported in the *in vivo* intestinal mucosa where subepithelial fibroblasts merge with other stromal cells in the lamina propria, for instance, interstitial fibroblasts and pericytes [48]. Besides not completely inadequate, this situation is not healthy to the cells. At D4, CCD18-Co cells continue to adopt an elongated shape with a slender cell body, but with a more mature and evolved shape (Fig. 4. 5 D).

After 21 days, 1×10^4 cells/cm² seems to be the healthier and well adapted cell density to a period time of 21 days (Fig. 4. 6 A), in detriment to the starry-like shape observed at higher densities (Fig. 4. 6 B-D). Until now, most of the assays were concordant in elect 1×10^4 cells/cm² as the cell density better adapted to 21 days of culture. To further characterize the 3D morphology of CCD18-Co and the surrounding matrix was adopted the lowest cell density of 1×10^4 cells/cm².

7th day

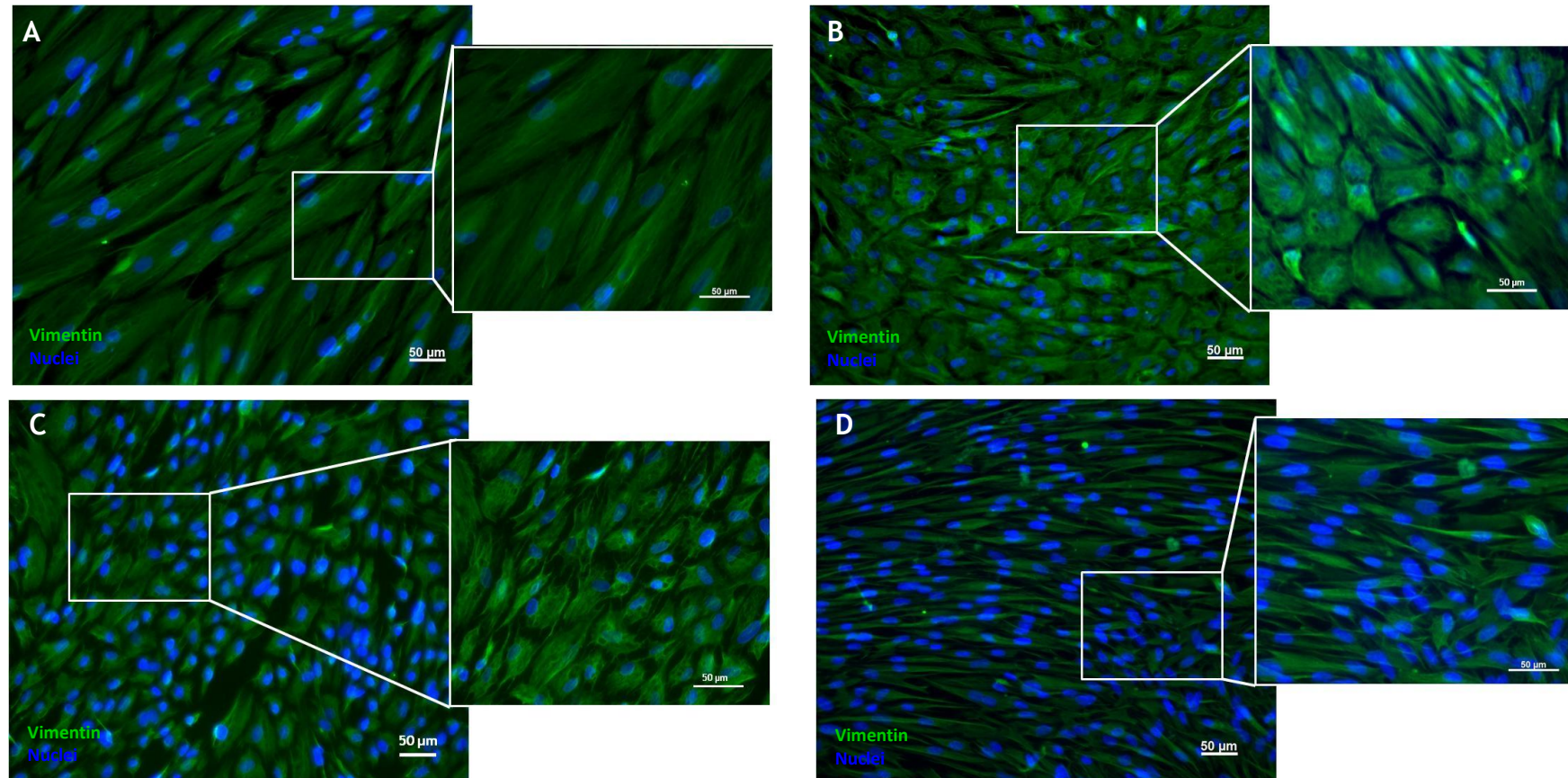


Figure 4. 4. Intestinal fibroblasts morphology identified by vimentin staining at day 7. After 7 days, are already notable differences in cells morphology. At D2 and D3, cells acquired a starry-like shape result of contact inhibition and lack of space. At D4, CCD18-Co cells presented skinny cell bodies noticing some cells overlapping. Vimentin was labelled with AlexaFluor488 (green) and a nucleus was labelled with DAPI (blue). Scale indicates 50 μm. (A) 1×10^4 cells/cm², (B) 5×10^4 cells/cm², (C) 10×10^4 cells/cm² and (D) 20×10^4 cells/cm².

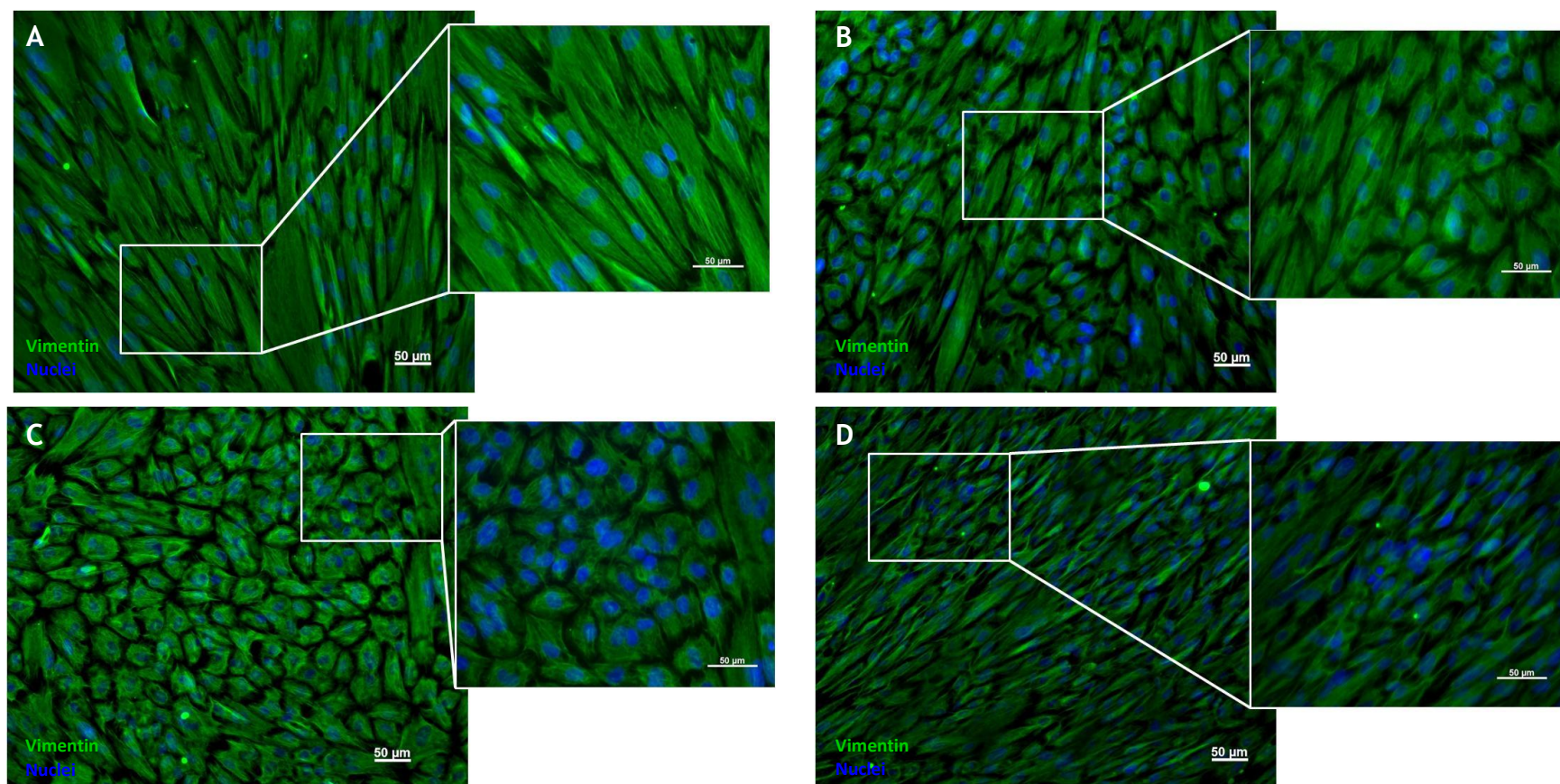
14th day

Figure 4. 5. Intestinal fibroblasts morphology identified by vimentin staining at day 14. The scenario is aggravated and, so far, D1 revealed the healthier cell density. The starry-like population is more prominent at higher densities (D2 and D3), observing the presence of multinucleated cells at D3. Vimentin was labelled with AlexaFluor488 (green) and a nucleus was labelled with DAPI (blue). Scale indicates 50 µm. (A) 1x10⁴ cells/cm², (B) 5x10⁴ cells/cm², (C) 10x10⁴ cells/cm² and (D) 20x10⁴ cells/cm².

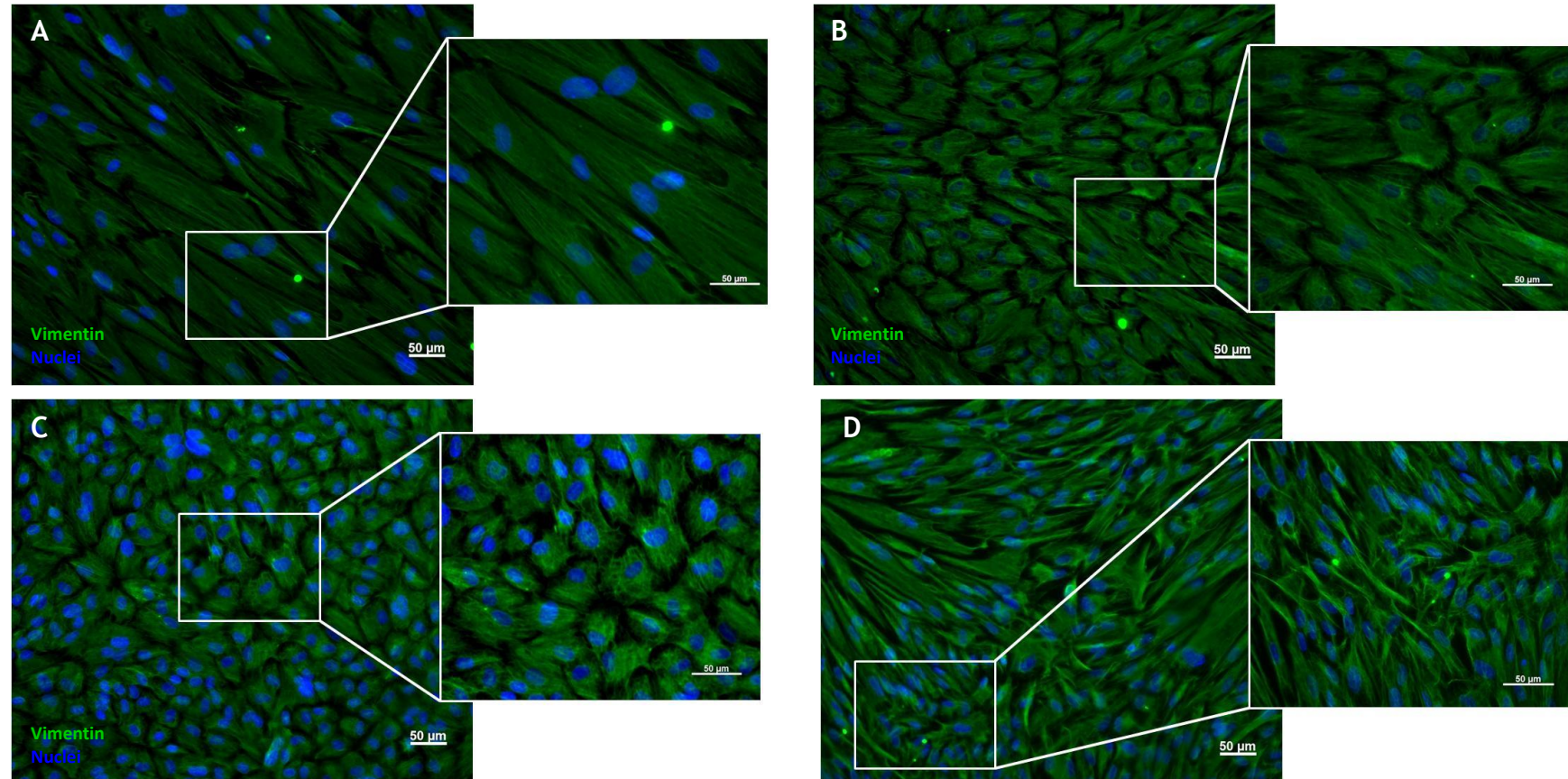
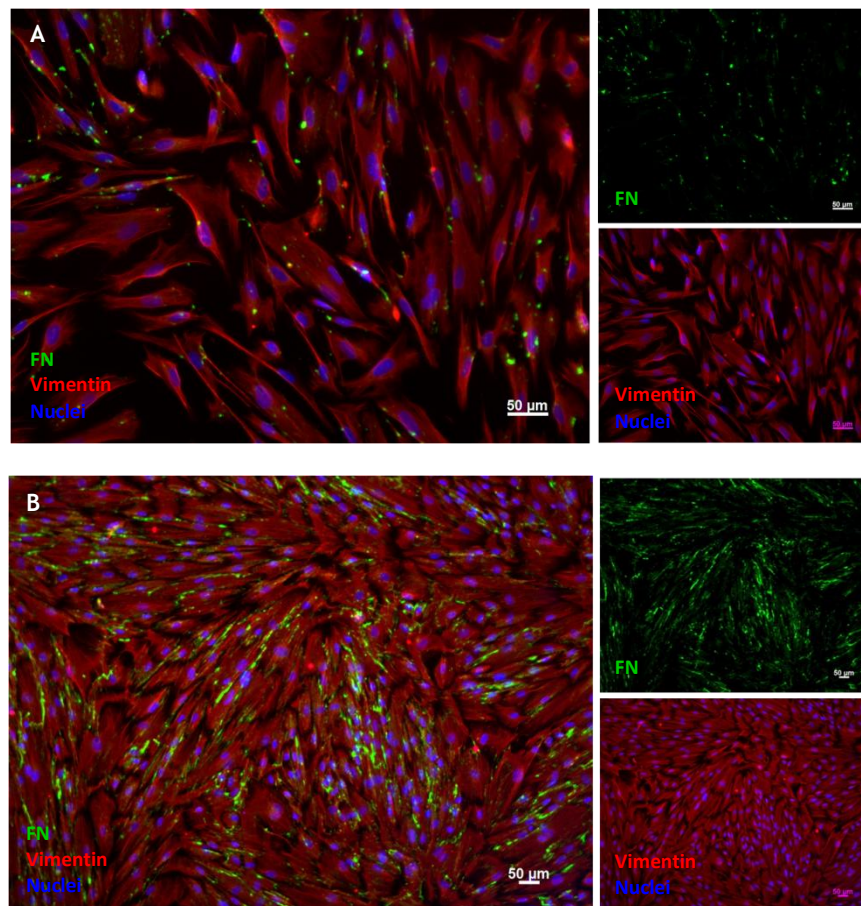
21st day

Figure 4. 6.Intestinal fibroblasts morphology identified by vimentin staining at day 21. After 21 days, D1 seems to be the well adapted density for 2D culture conditions observed by the distended cell shape and in detriment of the cell morphology at higher cell densities (D2, D3 and D4). Vimentin was labelled with AlexaFluor488 (green) and a nucleus was labelled with DAPI (blue). Scale indicates 50 μm. (A) 1x10⁴ cells/cm², (B) 5x10⁴ cells/cm², (C) 10x10⁴ cells/cm² and (D) 20x10⁴ cells/cm².

4.1.5. Characterization of ECM components

Fibroblasts signaling can substantially differ depending on whether the cells are in 3D or 2D culture conditions. The matrigel matrix affected cell metabolic activity and proliferation, as previously observed by resazurin assay and dsDNA quantification. Also, cell morphology was found to be susceptible to the 3D matrix constrictions. Additionally, fibroblasts are specialized cells in the remodeling of ECM and fibronectin (FN) is one of the components associated with biological processes involved in that remodeling [137]. FN characterization was thus performed over 21 days, simultaneously staining the intermediate filaments of CCD18-Co cells using vimentin to observe the changes in cell morphology in 3D conditions.

As can be observed in **Fig. 4. 7 A**, the FN staining after 7 days was condensed to small dots. After 14 days, the outcome was a more complex and concrete staining uniquely produced by CCD18-Co cells (**Fig. 4. 7 B**). At day 21, the levels of FN expression seemed not to be altered when compared with day 14 (**Fig. 4. 7 C**).



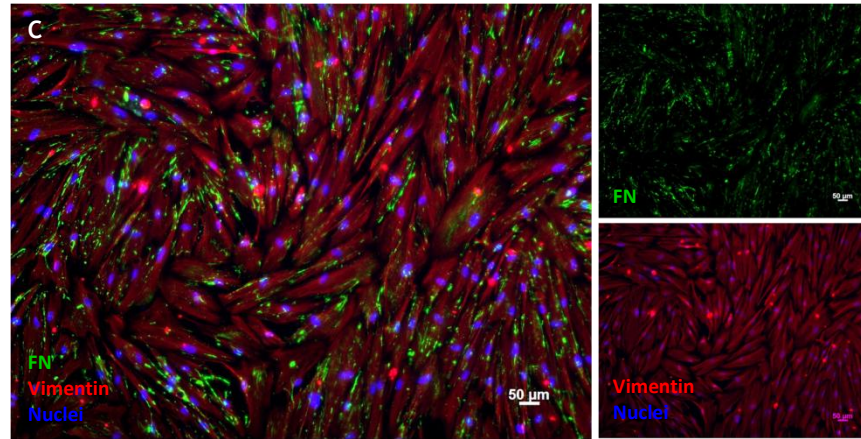


Figure 4. 7. Characterization of fibronectin network of CCD18-Co cells cultured in 2D standard conditions. Over 21 days, is observed a shy increase of FN expression. (A) At day 7 the FN is limited to condensed small dots, but after 14 days (B), it starts to reveal a more concrete staining. (C) At day 21, the levels of FN expression are maintained with similar complexity compared with day 14. Fibronectin was labelled with AlexaFluor 488 (green), vimentin AlexaFluor 594 (red) and nucleus with DAPI (blue). Scale indicates of 50 μm .

Regarding the FN expression in 3D conditions, was observed that after 7 days CCD18-Co cells revealed a complex network of FN (**Fig. 4. 8 A**). Even though manufactures report the detection of trace amounts of FN by Western Blot [139], matrigel is from rat origin [140] and FN expression was assessed with a human antibody to only assess the FN produced by CCD18-Co cells. Overall, matrigel seems to act as an initial livelihood that maintains the 3D rearrangement of fibroblasts while they are not producing their own ECM components. After 14 days, the FN expression was similar to the first 7 days (**Fig. 4. 8 B**). At this point, probably most of the existent matrigel was completely substituted by ECM components produced by CCD18-Co cells. To confirm this, further studies are required namely compare the expression of FN of rat origin and human origin. No relevant images were obtained for the 21 days.

Nonetheless, CCD18-Co cells showed to be evenly distributed throughout the matrigel matrix confirming that 1×10^4 cells/cm² was the cell density better adapted to both 2D and 3D culture conditions. Usually, there is no correlation between 2D and 3D mechanisms, but, in this specific case, it was established for cell density. As soon as matrigel polymerization was completed, matrigel thickness was significantly reduced. After 21 days under the effect of CCD18-Co cells and environmental factors, matrigel content was probably substituted in the totality. In fact, matrigel not only contains basement membrane components (laminins, collagen IV and heparin sulphate proteoglycans) but also matrix degrading enzymes (MMP-2 and MMP-9) and growth factors (EGF, bFGF, NGF, PDGF, TGF- β) [141]. Even though the rate of matrigel degradation is not reported, MMPs are vital to the degradation of both basement membrane and stromal matrices. Gelatinases MMP-2 and MMP-9 and transmembrane MMPs are critical mediators of basement membrane remodelling [142]. Therefore, even though the matrigel degradation rate is not reported, its integrity is compromised by plenty of environmental factors that will obey the system to react and produce ECM components.

In spite the cell density has been the same for 2D and 3D culture conditions, the cell morphology showed to be quite distinct. When CCD18-Co cells are cultured in a 3D Matrigel matrix (**Fig. 4. 8 A and B**), they appeared elongated but spindle-shaped. This characteristic morphology was already observed in cell-derived 3D matrices [137]. Compared with the cells seeded on a 2D surface at the same density (1×10^4 cells/cm²), these ones possesses a characteristic fan-shaped lamellae.

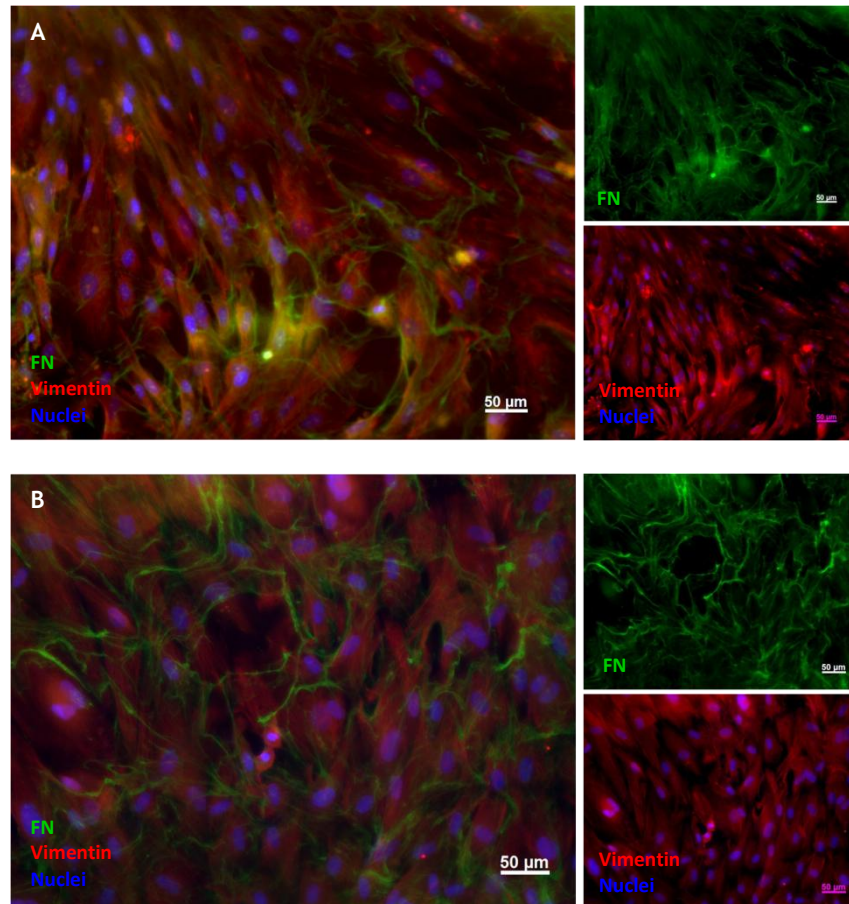


Figure 4. 8. Characterization of fibronectin network of CCD18-Co cells cultured in 3D conditions. (A) After 7 days CCD18-Co cells embedded in Matrigel® revealed a complex network of fibronectin comparable to day 14 (B). The 3D distribution of CCD18-Co cells seems well adapted to the matrix microenvironment. Fibronectin was labelled with AlexaFluor 488 (green), vimentin AlexaFluor 594 (red) and nucleus with DAPI (blue). Scale indicates 50 μ m.

Overall, CCD18-Co cells in 2D culture condition were broader and more flattened, presenting more cell protrusions than in the 3D matrices, tendency already observed by Hakkinen K. *et al.* [128].

4.1.6. Topographical features of CCD18-Co cells

The topographic features of the membrane were further visualized by SEM. After 21 days, CCD18-Co cells showed to populate the entire film as observed in **Fig. 4. 9 A**, however,

their elongated morphology do not fill all the spaces allowing a freer permeation. In the detached image of **Fig. 4. 9 A**, are represented the hole spaces between CCD18-Co cells acquired throughout the capture of retrodifused electrons instead of the standard secondary electrons. This acquisition delineates the regions of higher atomic number, appearing brighter [143]. In this case, the brighter regions represent the glass coverslip, translated in open spaces for freely permeation of a compound.

On the other hand, CCD18-Co cells seeded onto matrigel previously coated Transwell inserts revealed a complex network composed by ECM components (**Fig. 4. 9 B**) in contrast with fibroblasts seeded on the coverslip (**Fig. 20**). In certain regions was possible to observe the porous of the membrane since CCD18-Co cells are not completely opaque.

In one case or in the other, the disposition and morphology of CCD18-Co cells forms a cell monolayer with unoccupied spaces that allow a freer permeation.

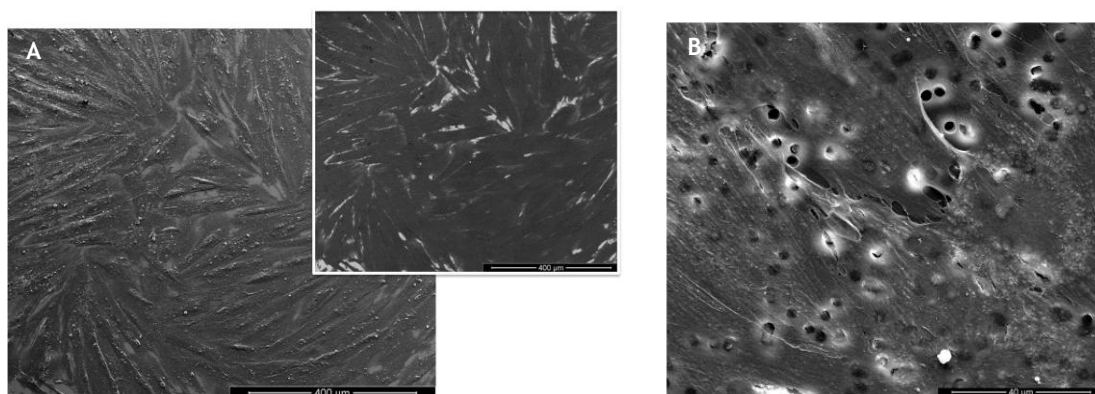


Figure 4. 9. Scanning electron micrographs of the CCD18-Co and Matrigel/CCD18-Co monolayer. (A) CCD18-Co monolayer in a coverslip showed to populate the entire film. Scale indicates 400 µm. (B) CCD18-Co cells seeded on Matrigel coated Transwell insert revealed a more complex network composed by ECM components. Scale indicates 40 µm.

4.2 Barrier modulation by epithelial cells

A characteristic feature of epithelial cells is the formation of cell junctions resulting in a tight cellular barrier that selectively controls the diffusion along the paracellular way, namely the TJs [144]. An intact barrier is crucial for the biological activities of a tissue but also for reliable *in vitro* experiments.

4.2.1 Assessing barrier integrity

A quantitative measure of integrity and maturity of a monolayer is the electric resistance of the cell layer [145]. Caco-2 cells are tighter linked by TJs, underestimating the paracellular permeability of compounds transported by this route. As result, Caco-2 cells are

characterized by high TEER values comprised between 150-1600 $\Omega \cdot \text{cm}^2$. This wide range is proof of the high inter-laboratory variability result of different culture conditions such as seeding density, passage number, days in culture, among others factors [87].

As observed in **Fig. 4. 10**, TEER measurements of the Caco-2 model presented higher values compared with the other models. In opposition, CCD18-Co model presented the lowest TEER values compared with the other models, what is justified by their morphological features that did not form a tight monolayer. The influence of matrigel coating with respect to the resistance values was not conclusive. Upon the introduction of CCD18-Co cells in direct contact with Caco-2 epithelial cells, TEER values presented a significant decrease. It may happen that CCD18-Co cells are interspaced between Caco-2 cells, interfering with the barrier tightness.

Were tested two distinct 3D configurations in which CCD18-Co cells are in different assemblies. As depicted in **Fig. 4. 10**, 3D co-cultures (CCD18-Co/Matrigel/Caco-2 and CCD18-Co+Matrigel/Caco-2) presented a slower rate of growth and significant lower TEER values over the 21 days when compared with 2D models. The CCD18-Co+Matrigel/Caco-2 model resulted in lower TEER values due to the fact that CCD18-Co cells dispersed in matrigel were not as proliferative as in 2D standard conditions, as observed before in the proliferative assays. Despite this, this model reached a desirable integrity above 200 $\Omega \cdot \text{cm}^2$ [146] and since this configuration closely mimics the cell-cell and cell-matrix interactions found in the intestinal mucosa, was proceeded with this configuration for further studies. Notwithstanding, the model CCD18-Co/Matrigel/Caco-2 represents an interesting configuration to study the *in vivo* deposition of ECM components by stromal that will form the supportive basement membrane [147].

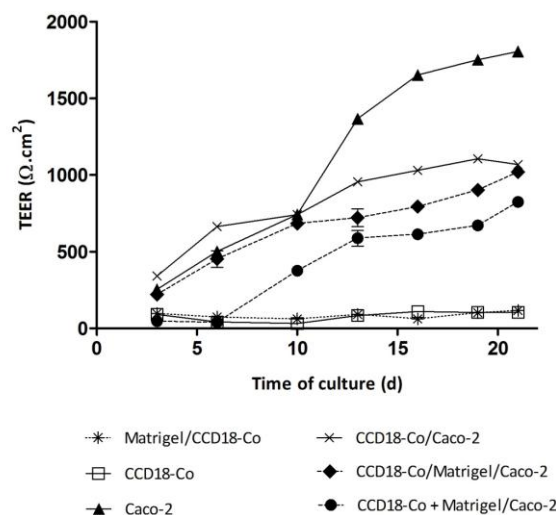


Figure 4. 10. TEER cell measurements monitored in function of time during 21 days of culture in cell based models of CCD18-Co intestinal fibroblasts and Caco-2 cells. Caco-2 monolayer was the model with higher TEER values in contrast with the CCD18-Co model. Results are the average of replicates and bars represent the standard deviation (n=4).

4.2.2. Epithelial-stromal interactions

In the sequence of the previous results, it was confirmed that the matrigel matrix influenced the metabolic activity and proliferation of CCD18-Co cells in 3D conditions.

To also clarify the interaction between stromal and epithelial cells, it was characterized the proliferative behaviour of CCD18-co cells and Caco-2 cells in direct contact after 7 days on a plastic substrate (Fig. 4. 11 A). Other control situations were tested as depicted in Fig. 4. 11 B - F.

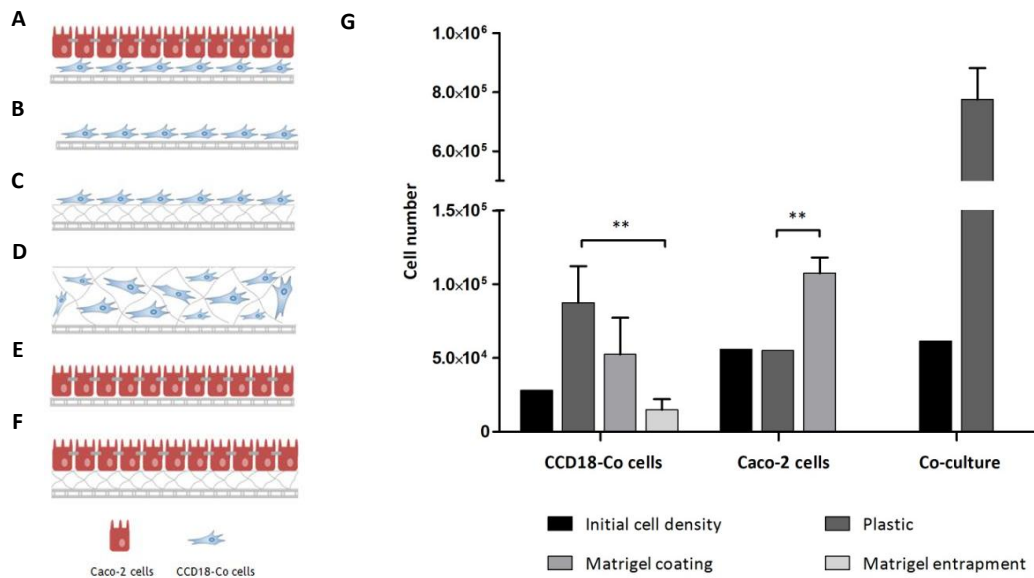


Figure 4. 11. Effect of the substrate on cell proliferation of Caco-2 and CCD18-Co cells after 7 days of culture. (A) Epithelial and stromal co-culture seeded onto a plastic substrate. (B) Stromal cell monoculture of CDD18-Co cells seeded on a plastic substrate. (C) Stromal monoculture of CCD18-Co cells seeded on a matrigel-coated plate. (D) Stromal cell monoculture of CDD18-Co cells embedded in a Matrigel® matrix. (E) Epithelial cell monoculture of Caco-2 seeded on plastic substrate. (F) Epithelial cell monoculture of Caco-2 seeded on a matrigel-coated plate. (G) CCD18-Co cells were less proliferative when entrapped in the Matrigel® matrix compared with plastic. In contrast, Caco-2 cells were significantly more proliferative when seeded onto a matrigel-coated substrate. Results are the average of replicates and bars represent the standard deviation (n=3). The statistical differences ($p < 0.05$) are denoted by (*).

After 7 days of growth, the cell number was determined and compared to the initial cell number. As observed in Fig. 4. 11 G, stromal cells were proliferative on plastic but showed reduced proliferation when embedded in matrigel. This reinforces that the entrapment of CCD18-Co cells in a matrigel matrix limits the cell proliferation, due to the matrix constraints and the hypoxia environment. Moreover, the process of entrapment requires multiple pipetting steps to suspend a cell suspension in the matrigel matrix and that may lead to some shear-induced damage. In contrast, CCD18-Co cells seeded onto a matrigel coated plate showed to be proliferative. In turn, Caco-2 cells showed identical proliferation when seeded onto a plastic surface with no significant differences comparing with the initial cell seeding, but significantly higher proliferation was observed when seeded onto matrigel

coated plates. These results were expected considering that matrigel is rich in ECM components that constitute an excellent substrate for cell growth and proliferation [148].

Taking advantage of the fact that CCD18-Co cells presented a proliferative behaviour in the plastic substrate, it was explored the effect CCD18-Co when in direct contact with Caco-2 cells. It was observed a high proliferative activity compared to the initial cell density, concluding that CCD18-Co cells did not inhibit the proliferation of Caco-2 cells and that both cell types had a synergetic effect when coated in direct contact.

4.2.3. Topographical features of Caco-2 cells

As observed in Fig. 4. 12, after 21 days, an uniform and differentiated character of Caco-2 epithelial cells was revealed by the presence of the typical microvilli, reported to be between 0.5-1.5 μm in length and 0.1 μm in width, varying according the species [149]. The boundary of cells is circumvented by a denser join of microvilli at the boundaries of two cells.

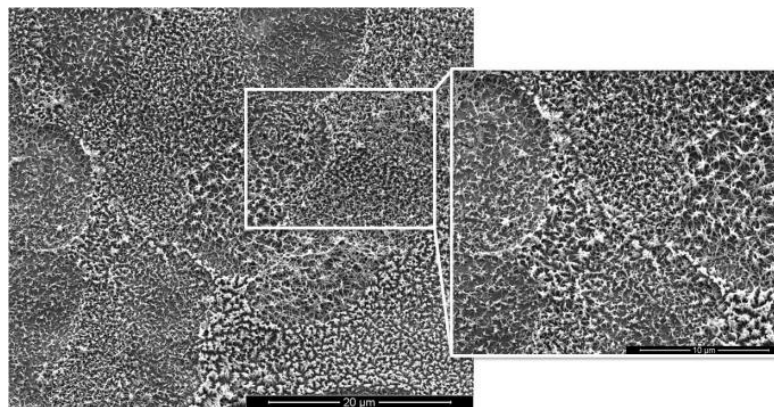


Figure 4. 12. Scanning electron micrographs of the Caco-2 monolayer after 21 days of culture. The presence of microvilli is the most characteristic feature of Caco-2 cells. Scale indicates 20 μm and 10 μm .

4.2.4. Expression of tight junctions proteins

Epithelial barrier function is regulated in large part by the apical intercellular junctions, referred to as the TJs, which comprises transmembrane proteins (occludin, claudin and junctional adhesion molecule (JAM)) and cytoplasmatic proteins (*zonula occludens* ZO-1, ZO-2, ZO-3 and cingulin) [150]. TJs form a continuous barrier between epithelial cells, which create tissue spaces and regulate the movement of solutes across the epithelium. The formation of a tight monolayer is a determinant factor to an *in vitro* model which application aims to test the absorption/permeation of a compound.

All the epithelial cells possess one common characteristic of express TJ proteins and form TJs between each other [144]. For instance, current evidences indicate that claudins play a central role in defining electrical resistance and permselectivity [151].

As observed in **Fig. 4. 13 A-C**, the top scanning of Caco-2 monolayer demonstrated a homogeneous ring along the perimeter of the cells confirming that Caco-2 cells formed tight and adherent junctions that limited the passage of components and confirm the functional properties of the barrier. This tendency had been already noticed by the TEER measurements, which returned high values of resistance.

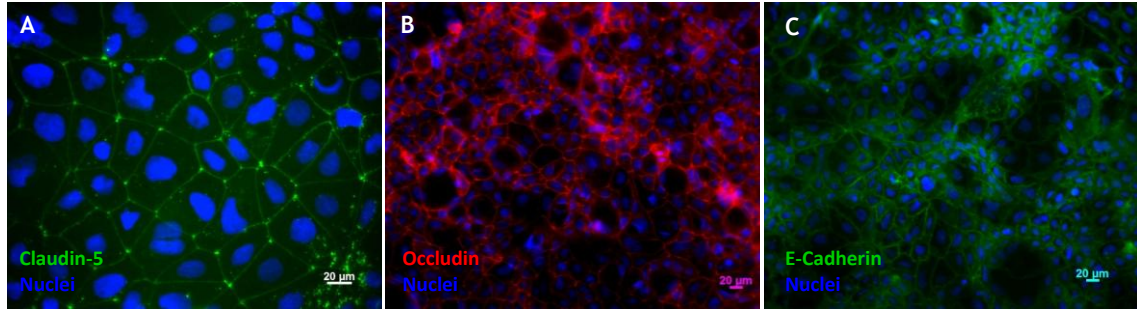


Figure 4. 13. Expression of claudin-5, occludin and E-cadherin in Caco-2 cell monolayer. (A) Claudin-5, (B) Occludin and (C) E-Cadherin. The top scanning of Caco-2 monolayer demonstrated a homogeneous ring along the perimeter of the cells, confirming the tight character of Caco-2 monolayer. Claudin-5 and E-Cadherin were labelled with AlexaFluor488 (green), occludin with AlexaFluor594 (red) and nucleus with DAPI (blue). Scale indicates 20 µm.

4.2.5. Caco-2 / HT29-MTX co-culture model

The introduction of new cell lines somehow interfere with the tighter character of the Caco-2 monolayer, but at the same time approximate the TEER values from the human intestine, ranging from 12-69 $\Omega \cdot \text{cm}^2$ [79, 102]. The introduction of HT29-MTX in the model interferes with the barrier formation as these cells will be interspersed between Caco-2 cells altering the tighter character of the Caco-2 monolayer. Even though both cells types establish TJ between each other, they present a structural variability facilitating the passage of ions by paracellular pathway [144].

The less tight character of the co-culture of Caco-2 and HT29-MTX cells was confirmed by the measurement of the TEER (**Fig. 4. 14**). The Caco-2/HT29-MTX monolayer presented a gradual increase of TEER values during 21 days but significantly lower when compared with the Caco-2 monolayer. The 3D triple model (CCD18-Co+Matrigel/Caco-2/HT29-MTX cells) revealed a decrease in TEER values compared with the previous 3D double model (CCD18-Co+Matrigel/Caco-2) result of the introduction of HT29-MTX cells.

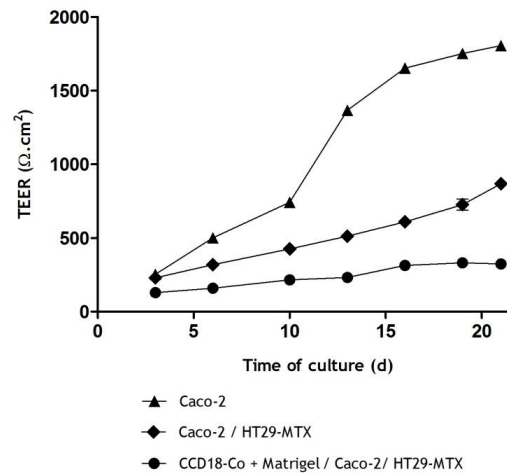


Figure 4. 14. TEER cell measurements monitored in function of time during 21 days of culture in a cell based model of CCD18-Co intestinal fibroblasts, Caco-2 and HT29-MTX cells. Caco-2/HT29-MTX model presented lower TEER values compared with Caco-2 model regarding the less tight character of tight junctions between both cell types. Results are the average of replicates and bars represent the standard deviation (n=4).

In Fig. 4. 15 A, it is depicted the expression of a tight junction protein claudin-5 between HT29-MTX cells, observing the rounder shape distinct from Caco-2 cells. When in co-cultured, Caco-2 cells and HT29-MTX cells also express a tight junction protein occludin (Fig. 4. 15 B), being impossible to discriminate Caco-2 cells from HT29-MTX by similar shape and size [77].

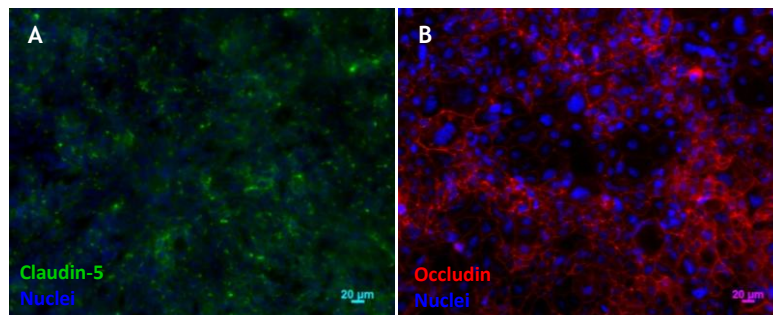


Figure 4. 15. Expression of claudin-5 in HT29-MTX monolayer and occludin in Caco-2/HT29-MTX monolayer. Despite the different structure of tight junctions of HT29-MTX cells, Caco-2/HT29-MTX monolayer expressed a strong staining for occludin tight junction protein. Claudin-5 was labelled with AlexaFluor488 (green), occludin with AlexaFluor594 (red) and nucleus with DAPI (blue). Scale indicates 20 μm.

In contrast, it was possible to distinct Caco-2 from HT29-MTX cells by transmission electron microscopy. As observed in Fig. 4. 16 A, the monolayer is richly covered by mucin-filled granules, which tend to coalesce, a process in which two or more elements merge to form a single element. The Caco-2 epithelial cells formed microvilli on their apical surface exhibiting slight variability in density along the different regions of the monolayer. HT29-MTX cells were distinguished by the presence of intracellular granules, appearance similar to *in vivo* goblet cells that contain mucin granules located below the apical membrane [152]. As depicted in Fig. 4. 16 B, both epithelial Caco-2 and HT29-MTX cells were able to form tight junctions.

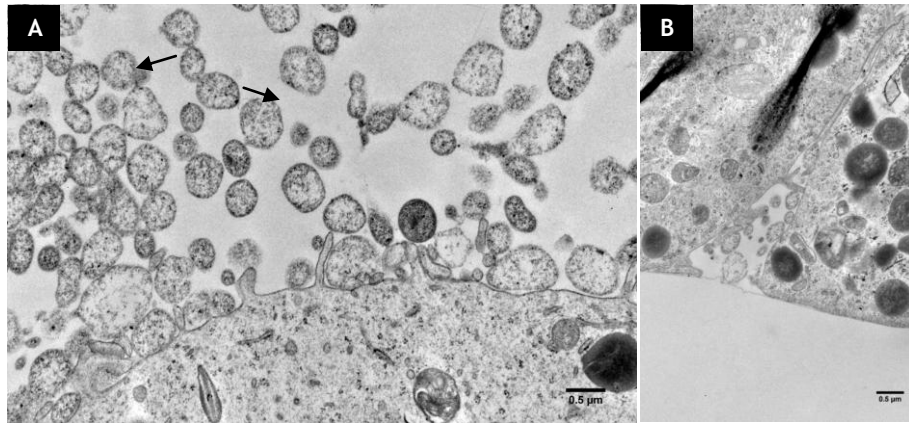


Figure 4. 16. Transmission electron micrographs of Caco-2/HT29-MTX monolayer. (A) Caco-2/HT29-MTX monolayer with typical microvilli and mucin granules. Scale bar indicates 0.5 µm. The black arrows represent indications of coalescence. (B) Tight junction formed between Caco-2 (left) and HT29-MTX cell (right). Scale indicates 0.5 µm. The fold is an artefact of specimen preparation procedure for TEM.

4.2.6. Evidence of M-like cell formation

M cells, located in the epithelia overlying mucosa-associated lymphoid tissues (i.e. PPs), are an instrument in initiating mucosal immunity and a portal for the delivery of macromolecules and microorganisms, consequently explored to antigen delivery [76]. The little known about the interactions between M cells and enterocytes and between M cells makes difficult their unambiguous identification. Claudin-1/3 and ZO-1 were detected in the TJs of both M cells and enterocytes within mouse intestinal follicles-associated epithelium (FAE) revealing the contact of M cells with the adjacent cells through TJs. Regarding AJs between M cells of mouse PPs, they revealed enhanced expression of β -catenin, α -actinin, polymerized actin and in some areas, E-cadherin, differing from enterocytes [153]. Furthermore, M cells appear to exhibit an increased depth and an altered rearrangement of TJ strands differing from the non-FAE intestinal epithelia [154].

TJs rearrangement between enterocytes and M cells is somehow reproduced *in vitro* by the conversion of Caco-2 epithelial cells into M-like cells by Raji B lymphocytes [36]. It was expected that the TEER values of Caco-2/Raji B model presented similar values to the Caco-2 model since this model is mainly comprised by Caco-2 cells. However, it was observed a decrease in TEER values compared with Caco-2 model that can be result of the conversion of Caco-2 cells into M-like cells. This seems to be a remote idea since M-like cells were originated from Caco-2 cells, which upon the introduction of Raji B cells (day 15), already form a very tight monolayer. Even though this evidence was already reported [106], is necessary to confirm this assumption by identifying the soluble factors responsible for this phenotypic induction and TJ strength.

The triple model of Caco-2/HT29-MTX/Raji B cells presented lower TEER values when compared with the Caco-2/Raji B control result of the presence of HT29-MTX cells. The 3D quadruple model (CCD18-Co+Matrigel/Caco-2/HT29-MTX/Raji B) presented a slower increase

of TEER values with a similar profile to the 3D triple model (CCD18-Co+Matrigel/Caco-2/HT29-MTX).

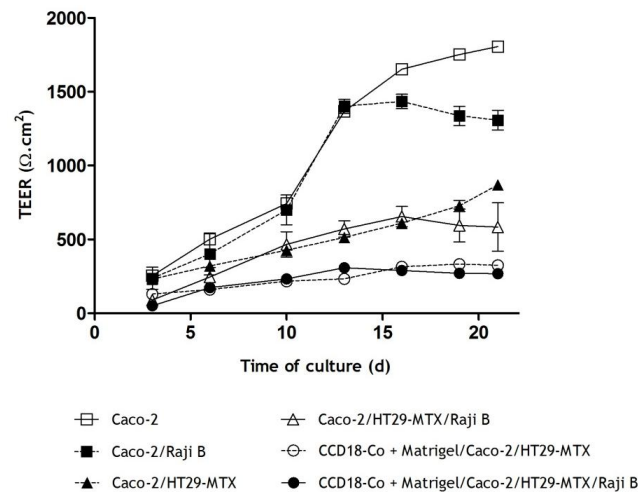


Figure 4. 17. TEER cell measurements monitored in function of time during 21 days of culture in a cell based model of CCD18-Co intestinal fibroblasts, Caco-2 cells, HT29-MTX and Raji B cells. Caco-2/Raji B model presented high TEER values, in the same range of the TEER values of Caco-2 model. Results are the average of replicates and bars represent the standard deviation (n=4).

In Fig. 4. 18, is possible to visualize that the triple model of Caco-2, HT29-MTX and M-like cells is tight enough despite the structural differences between the epithelial cells.. The top scanning of co-culture monolayer demonstrates a homogeneous ring around the perimeter of the cells, morphologically identical despite the heterogeneous cell population.

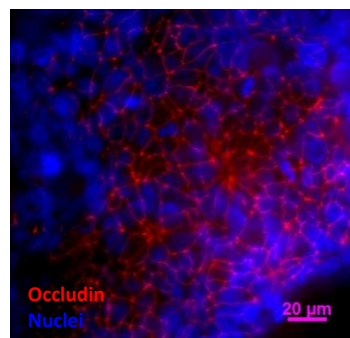


Figure 4. 18. Expression of tight junction protein occludin in Caco-2/HT29-MTX/Raji B model. Despite the structural differences between epithelial cells, the triple model revealed a tight character. Occludin was labelled with AlexaFluor594 (red) and a nucleus was labelled with DAPI (blue). Scale indicates 20 μm.

4.2.6.1. Quantification of alkaline phosphatase activity

Morphologically distinct from enterocytes, M cells display poorly organized brush border with short irregular microvilli at the apical domain. As result, the enzymatic activity of alkaline phosphatase is down-regulated [8]. *In vitro* studies showed that lymphocytes trigger the conversion of enterocytes cell lines into M-like cells in co-culture under influence of

membrane-bound lymphotoxin (LT α 1B2) secreted by B-lymphocytes [36]. Therefore, the presence of Raji B lymphocytes in the basolateral side is known to protagonize the M-cell formation [106-108].

Alkaline phosphatase (ALP) is as an excellent marker of crypt-villus differentiation, since ALP of Caco-2 cells have identical sequence [155] and similar biochemical characteristics [156] compared to the intestinal alkaline phosphatase (IAP) isoenzyme, which is exclusively expressed in villi enterocytes [157].

In order to trigger M-like cell formation, an alkaline phosphatase assay based on a work of Ferruza et al. [123], was performed as mean to monitor the down-regulation in the alkaline phosphatase activity. As observed in Fig. 4. 19, the ALP activity increased over the 21 days of culture, although not significantly between days 14 and 21 (period of contact with Raji B cells), what reveals that these assay was not significantly sensitive enough to trigger the presence of M-like cells. The gradual increase of ALP activity was already been observed in Caco-2 monolayer [123, 158]. Nonetheless, this quantitative assay was never applied to study the down-regulation of ALP in M-like cells and the down-regulation of ALP activity was only observed in *in vivo* studies. There are qualitative evidences of the decrease of ALP activity in a triple co-culture of Caco-2/HT29-MTX/Raji B when compared with the control of Caco-2/HT29-MTX [108], however further evidences are required. Moreover, it was not studied the amount of M-like cells formed regarding the initial Raji B cell density. This is certainly a hypothesis to explore, in order to equalize the *in vitro* cell number of M-like cells to the *in vivo* proportions of M cells.

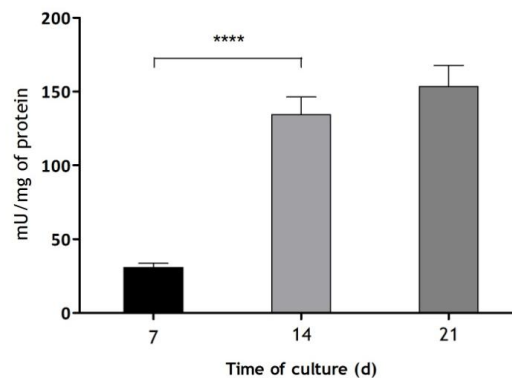


Figure 4. 19. Alkaline phosphatase (ALP) activity during differentiation of Caco-2/Raji B model. ALP activity increased over the 21 days of culture revealing that this method was not sensitive enough to detect the presence of M-like cells. Results are the average of replicates and bars represent the standard deviation (n=3). The statistical differences ($p < 0.05$) are denoted by (*).

4.2.6.2. Morphological features of M-like cells

HT29-MTX cells display more sparse microvilli [77], what could enter in conflict with the identification of M-like cells. Opportunely, HT29-MTX cells contain intracellular mucin granules that allowed the identification of HT29-MTX cell by transmission electron

microscopy. As observed in **Fig. 4. 20 A**, after 21 days some HT29-MTX cells are found closer to the basal side of Caco-2 cells. In fact, HT29-MTX present a doubling time different from Caco-2 cells, the last with higher doubling time [106]. Notwithstanding, are still observed mucin-filled granules at the epithelial monolayer (**Fig. 4. 20 B**) which shows that, although not pictured, some HT29-MTX cells are near the apical side of Caco-2 cells, releasing some mucin granules that remain adsorbed to the apical membrane. Hilgendorf *et al.* had already noticed that the standard composition of Caco-2 cell medium did not favored the HT29-MTX growing, leading to co-cultures that were more similar to Caco-2 monocultures than mixed cultures [105].

Once again, Caco-2 cells displayed slight variability in microvilli density along of the different regions of the monolayer. This variability did not allow the undoubted confirmation of the presence of M-like cells. Nonetheless, epithelial cells are tightly linked each other constituting an evident barrier to the drug translocation (**Fig. 4. 20 B**).

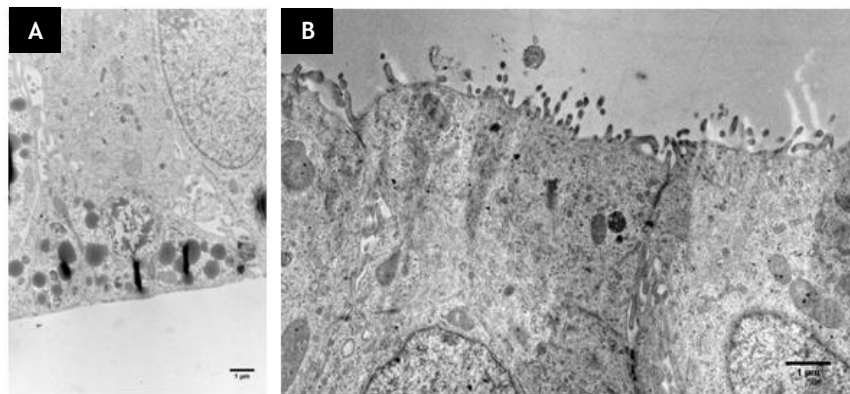


Figure 4. 20. Transmission electron micrographs of Caco-2/HT29-MTX/Raji B monolayer. (A) HT29-MTX cells with intracellular mucin-filled granules sited over the Transwell filter. Scale indicates 1 µm. (B) Tight junctions formed between epithelial cells with typical microvilli and some mucin granules. Scale indicates 1 µm. The fold is an artefact of specimen preparation procedure for TEM.

Nevertheless, M-like cells were possibility identified by scanning electron microscopy by the lack of microvilli interspersed between Caco-2 cells (**Fig. 4. 21 A**). It was not possible to distinguish Caco-2 cells from HT29-MTX cells by their morphological features but residual evidences of mucus seem to be retained over the surface of microvilli (**Fig. 4. 21 B**). However, this did not preclude the presence of an uninterrupted layer of mucus covering the epithelial layer, as reported by other authors in the same proportions of Caco-2 and HT29-MTX cells [118]. Probably, most of the mucus was lost during the membrane processing for scanning electron microscopy.

Overall, SEM allowed to observe a more representative area to the identification of M-like cells while TEM was more effective in the identification of HT29-MTX cells and presence of mucin-filled granules.

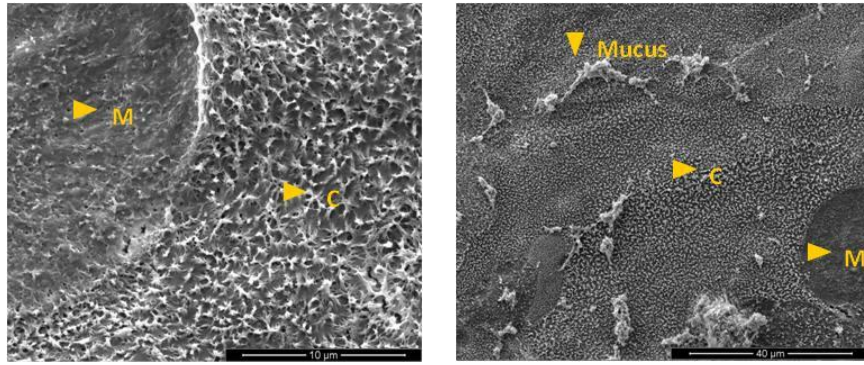


Figure 4. 21. Scanning electron micrographs of the Caco-2/Raji B double model and of the Caco-2/HT29-MTX/Raji B triple model after 21 days. (A) M-like cells are identified by shorter or lack of microvilli in the Caco-2/Raji B model. (B) HT29-MTX cells were not distinguished from Caco-2 cells but there are evidences of mucus attached. The cracking is an artefact of specimen preparation procedure for SEM. Scale bar of 50 µm and 40 µm.

4.2.7. Visualization of the tridimensional model

As observed in **Fig. 4. 22 A**, Caco-2 cells presented their characteristic columnar shape. After three weeks of culture, a dense monolayer of epithelial cells was maintained over a gel layer of CCD18-Co fibroblasts embedded in matrigel (**Fig. 4. 22 B**). This organization is somewhat like the native small intestinal mucosa in with the epithelial cells lying onto a lamina propria (**Fig. 4. 22 C**) [111]. In fact, the initial thickness of the matrigel was lost but their degradation allowed the CCD18-Co cells to easier populate the 3D matrix. Therefore, CCD18-Co cells had an active role in the construction of lamina propria that sustains epithelial cells. The elongated morphology of CCD18-Co cells in a 3D matrix seems to indicate that CCD18-Co cells were able to modulate the matrix, otherwise would appear less elongated.

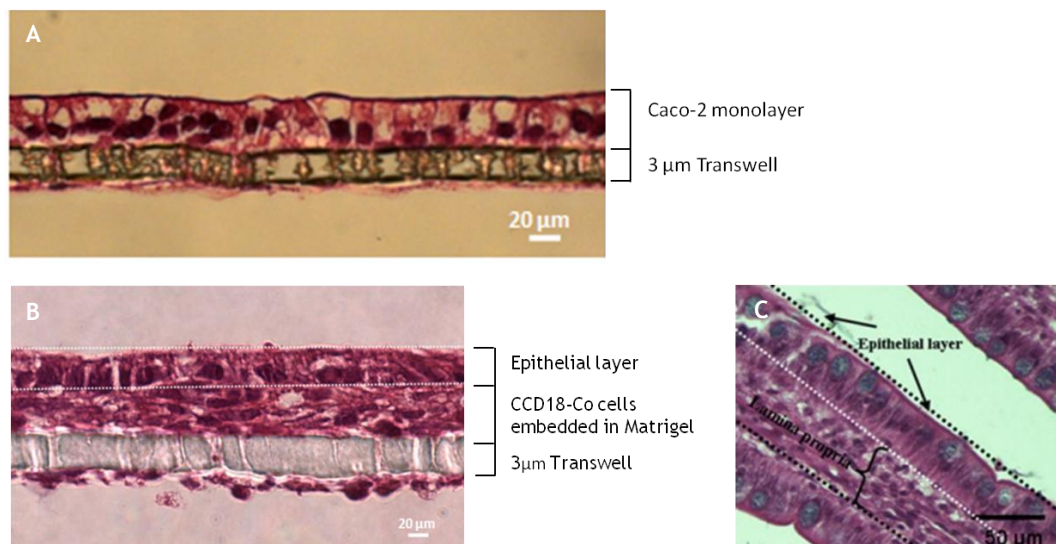


Figure 4. 22. Tridimensional model visualization through histological cut stained with H&E. (A) Caco-2 monolayer showing the characteristic columnar shape of Caco-2 cells. The colouration below the Transwell insert is an artefact of the histological staining. (B) In vivo cut of the in vivo small intestinal mucosa. (C) CCD18-Co intestinal fibroblasts embedded in matrigel sustain Caco-2 cells over them. Scale indicates 20 µm.

Then, the matrix that sustains the epithelial cells was characterized by staining fibronectin. As observed in Fig. 4. 23, the FN expression was restricted to the fibroblastic layer that sustains the Caco-2 monolayer. Some fibroblasts were able to cross the 3 μ m Transwell insert, evidenced by the presence of residual staining above the filter. This result is an evidence of the active role of CCD18-Co cells, as they were able of reformulate the matrigel and leave the matrix.

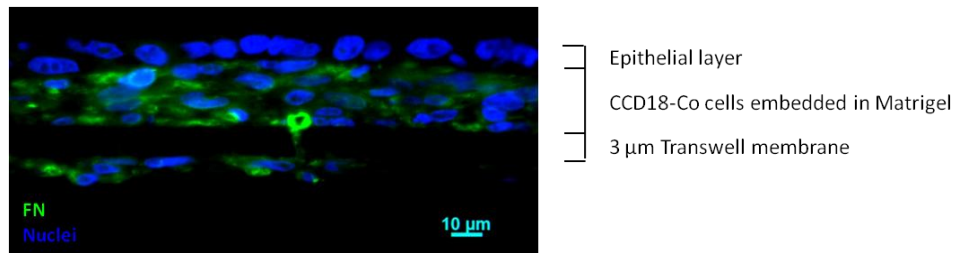


Figure 4. 23. Fibronectin expression of a paraffin section of tridimensional model of CCD18-Co cells embedded in Matrigel and Caco-2 cells. FN is restricted to the fibroblastic layer of CCD18-Co cells demonstrating the supportive role of CCD18-Co cells in maintain epithelial cell layer. Fibronectin was labelled with AlexaFluor 488 (green) and nucleus with DAPI (blue). Scale indicates 10 μ m.

4.2.8. Insulin permeability

The functional character of the model was validated comparing the permeability profile of insulin in the monocultures and co-cultures of CCD18-Co, Caco-2, HT29-MTX and Raji B cells. The choice of insulin as a model drug supported by the previous experience of the group.

Besides the controversial character of the absorption mechanism of oral administrated insulin, the paracellular pathway is the most feasible pathway for its passage across the intestinal epithelium due to its high hydrophilicity [15]. Moreover, since insulin (isoelectric point \approx 5.4) is negatively charged at the pH of the small intestine, its interaction with negatively charged mucins at the epithelial surface will repulse insulin avoiding it to reach the cells [159, 160]. In contrast, morpho-cytochemical and biochemical evidences in both normal and diabetic rats studied the ileal tissues and between 2 and 5 minutes, insulin labeling was associated to the microvilli and endocytic vesicles in the apical portion of epithelial cells. At 10 minutes, insulin labeling was found at the level of the trans-side of Golgi apparatus and along the basolateral membranes of epithelial cells [161, 162]. Importantly, insulin has been reported to be absorbed into the rat ileal epithelium in the presence of permeation enhancers and protease inhibitors [162], but in addition, an insulin-degrading enzyme (IDE) has been identified in the cytosol of the rat intestinal enterocytes constituting up to 92% of total insulin-degrading activity [163]. Moreover, insulin receptors, that have been identified in the basolateral membranes of dog intestinal mucosa [164], in mouse intestinal cells [165] and in the membrane of Caco-2 cells [166], may be functional in the protein transport across the intestinal epithelium but their precise role remains unclear.

Thus, both pathways are feasible to transport insulin. However, the endosomal degradation and the insulin-degrading enzyme neglect the real contribution of the transcellular pathway in the transport of insulin.

Permeability studies were performed with insulin in free-form over 4 hours with parallel measurement of TEER values. Before the experiment, the cultures were equilibrated with HBSS and TEER values were allowed to stabilize. The intrinsic resistance of the cell layer is influenced by the cell culture medium in the apical and basolateral compartment, by the membrane of the filter inserts and by the electrode-medium interface. Since transport experiments are performed in HBSS, TEER values typically decreased in result of the adjustment of the experiment conditions.

As depicted in Fig. 4. 24, all the models showed the same profile in the TEER measurement, reaching an almost stationary profile after 1 hour of experiment. As the compound interacts with the cell layer, TEER values usually decrease resulting in the opening of TJs. That decrease is returned as soon as the culture conditions are reestablished as proved by higher values of FITC-Dextran permeation and reported by other authors [167, 168].

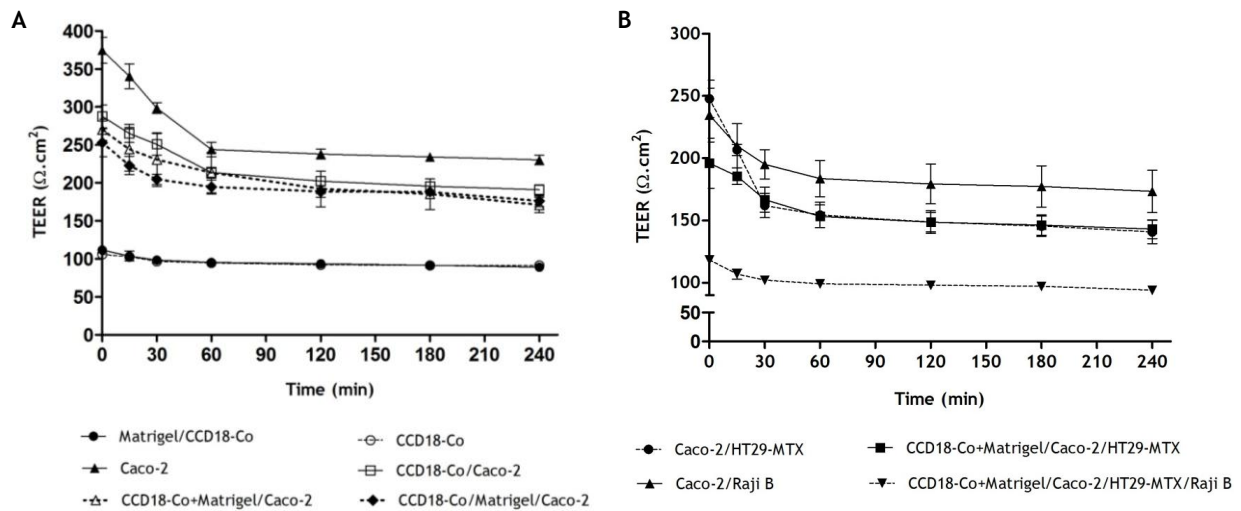


Figure 4. 24. TEER values monitored in function of time during 4 hours of permeability experiments, comprising Caco-2, CCD18-Co cells, HT29-MTX and Raji B cells. All models revealed a significant decrease in TEER values. Despite the decrease, Caco-2 model presented the higher TEER values after 4 hours. Results are the average of replicates and bars represent the standard deviation (n=3).

4.2.8.1. Insulin quantification

The permeability profile of all the models follows a similar tendency during the time of the experiment translated in an increased permeability with a decrease of TEER values (Fig. 4. 25).

Caco-2 monoculture presented the lower amount of permeated insulin, expected by the higher TEER values and the tighter junctions. Noticeably, as higher the initial TEER values,

the profile of decrease of resistance values during the permeability experiment seems to decrease quicker and more accentually, with exception of the monolayer co-culture of CCD18-Co and Caco-2 cells. In contrast, CCD18-Co and Matrigel/CCD18-Co monolayer allowed the higher permeation to the passage of insulin without meaningful variations after 30 minutes (Fig. 4. 25). The effect of matrigel coating is not perceived in TEER values, but becomes more noticeable that matrigel coating seems to interfere with the permeability studies, limiting the transport of insulin to the basolateral side. This observation is justified by the proliferative behaviour of CCD18-Co cells when seeded on a matrigel substrate. The permeability profile of the CCD18-Co/Caco-2 model was similar to the Caco-2 model, since both cells co-exist in co-culture as observed by their proliferative character.

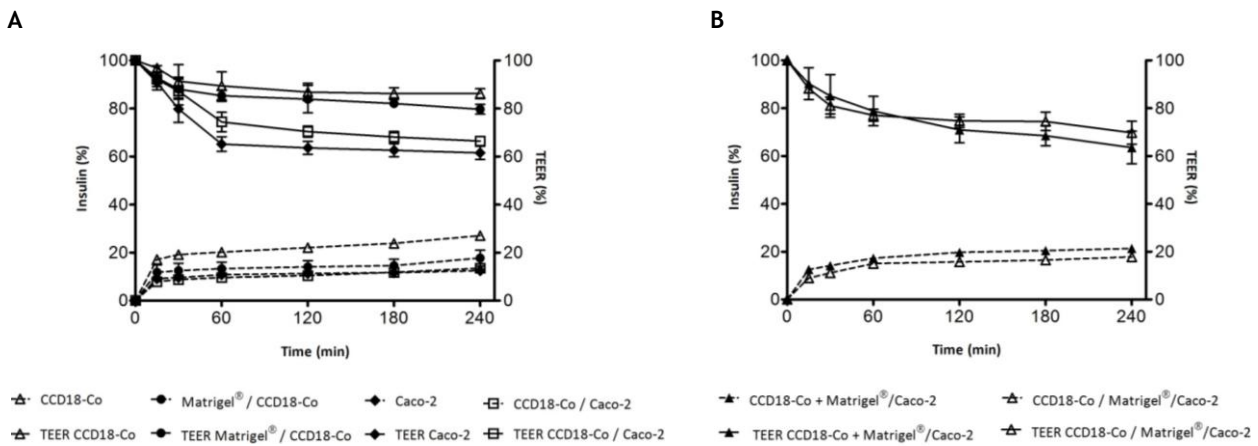


Figure 4. 25. Cumulative transport of insulin across monocultures and co-cultures of CCD18-Co and Caco-2 cells. CCD18-Co model is the one presenting higher permeation to insulin regarding the high pore size of CCD18-Co monolayer. Results are the mean of replicates and bars represent the standard deviation (n=3).

The apparent permeability coefficient (P_{app}) represents a cumulative amount of the test compound in the basolateral side, not distinguishing the contribution of each transport pathway. The P_{app} value to insulin in Caco-2 monolayers was higher 1×10^{-6} cm/s but, as expected, presented the lower P_{app} compared with the other models (Table 4. 1). Moreover, P_{app} values of CCD18-Co/Caco-2 model were not significantly different of Caco-2 model, recalling the similarity of TEER values with the Caco-2 model. CCD18-Co cell seeded onto matrigel coating presented significantly lower P_{app} compared with the P_{app} of CCD18-Co monolayer. As observed before by scanning electron microscopy, CCD18-Co seeded onto matrigel presented a complex matrix that can offer more resistance to the transport of insulin (Fig. 4.9 B). Comparing both 3D configurations, the 3D double model CCD18-Co+Matrigel/Caco-2 allowed a higher permeation of insulin when compared with the CCD18-Co/Matrigel/Caco-2. Besides the less proliferative character of CCD18-Co cells when embedded in matrigel, the 3D assembly seems to be a critical factor in the transport of insulin.

Table 4. 1. Apparent permeability coefficient ($P_{app} \times 10^{-6}$ cm/s) of insulin across the monolayer comprising Caco-2 and CCD18-Co cells. Results are the mean of three replicates \pm standard deviation ($n=3$, mean \pm SD).

<i>In vitro</i> models	P_{app} ($\times 10^{-6}$ cm/s)	Significance level
CCD18-Co	6.03 ± 0.23	(***)
Matrigel/CCD18-Co	3.95 ± 0.74	
Caco-2	2.77 ± 0.18	(ns)
CCD18-Co/Caco-2	3.00 ± 0.08	
CCD18-Co +Matrigel/Caco-2	4.76 ± 0.08	(**)
CCD18-Co/Matrigel/Caco-2	3.96 ± 0.19	

As stated before, the introduction of mucus-producing cells interferes with the barrier tightness. These results showed that HT29-MTX cells were involved in the increase of permeability of insulin (Fig. 4. 26). In the *in vivo* scenario, mucus interaction with the negative charges of insulin difficult their entrapment by the mucus, however, in static conditions the mucus may approximate the insulin molecules near of the epithelial surface, retaining a considerable amount that will permeate through the less strait TJ between both epithelial Caco-2 and HT29-MTX cell lines. Therefore, the existence of mucus was propitious to the transport of insulin, effect already reported by Jin *et al.* [169].

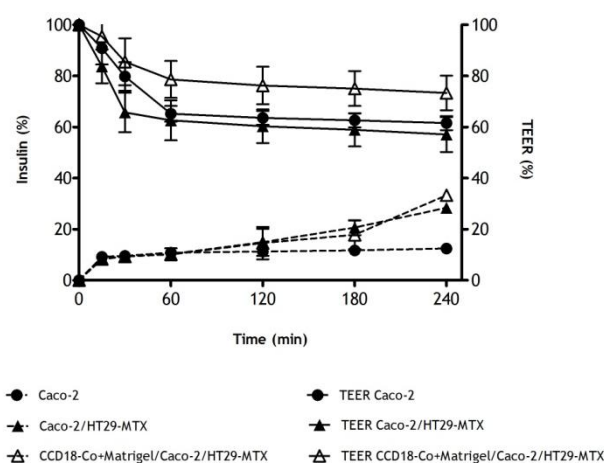


Figure 4. 26. Cumulative transport of insulin across monocultures and co-cultures of CCD18-Co fibroblasts, Caco-2 cells and HT29-MTX cells. Caco-2/HT29 model presented higher permeation to insulin compared to the Caco-2 model control, revealing that the existence of mucus was propitious to the transport of insulin. Results are the mean of three replicates and bars represent the standard deviation ($n=3$, mean \pm SD).

Comparing the Caco-2/HT29-MTX model with the Caco-2 control, the double model presented higher susceptibility to the permeation of insulin favored by the presence of mucus and the less tight junctions between both Caco-2 and HT29-MTX epithelial cells. As expected, the 3D triple model (CCD18-Co+Matrigel/Caco-2/HT29-MTX) showed relative higher

permeation compared with the 3D double model (CCD18-Co+Matrigel/Caco-2) result of the presence of HT29-MTX cells (Table 4. 2). As argued before, the 3D models allowed a higher permeation of insulin justified by entrapment of CCD18-Co cells in the matrigel matrix.

Table 4. 2. Apparent permeability coefficient ($P_{app} \times 10^{-6}$ cm/s) of insulin across the models comprising CCD18-Co, Caco-2 and HT29-MTX cells. Results are the mean of three replicates \pm standard deviation ($n=3$, mean \pm SD).

<i>In vitro</i> models	P_{app} ($\times 10^{-6}$ cm/s)	Significance level
Caco-2	2.77 ± 0.18	***
Caco-2/HT29-MTX	6.32 ± 0.18	
CCD18-Co + Matrigel/Caco-2	4.76 ± 0.08	**
CCD18-Co + Matrigel/Caco-2/HT29-MTX	7.42 ± 0.28	

M cells are characterized by their high transcytotic capacity and reduced expression of enzymatic activity throughout the down-regulation of alkaline-phosphatase and sucrase enzymes present in microvilli [8]. Even though the *in vitro* presence of M-like cells has been progressively revealed [106, 108], detailed information lacks regarding the permeability behaviour of the models comprising these cells.

In the results depicted in Fig. 4. 27, seems that the presence of M-like cells increased the permeability of insulin. In fact, it remains the question whether, upon Raji B cells conversion, there is any change in TJ structure that would justify an increase in permeability. If there is a loss of the tightness of the monolayer due to the presence of M-like cells, that would favor the paracellular route. More remotely, insulin can also take advantage of the transcytotic capacity of M-like cells.

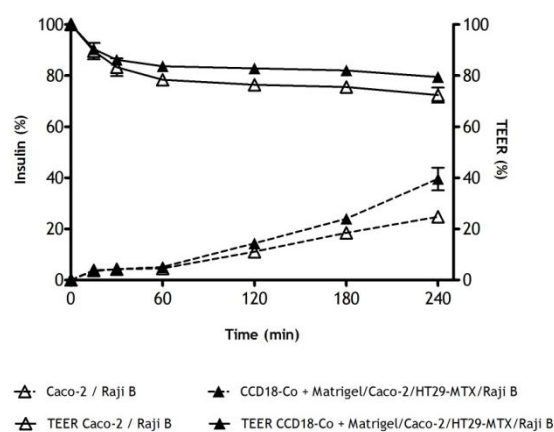


Figure 4. 27. Cumulative transport of insulin across monocultures and co-cultures of CCD18-Co fibroblasts, Caco-2 cells, HT29-MTX and Raji B cells. (A) Caco-2/Raji B model presented lower permeation to insulin compared to the CCD18-Co+Matrigel[®]/Caco-2/HT29-MTX/Raji B model (B), mainly result of the introduction of HT29-MTX cells. Results are the mean of three replicates and bars represent the standard deviation ($n=3$, mean \pm SD).

Comparing the P_{app} values, it is evident that the presence of M-like cells favored the transport of insulin. The 3D quadruple model (CCD18-Co+Matrigel/Caco-2/HT29-MTX/Raji B cells) represents the model that allows the more freely permeation of insulin (Table 4. 3).

Table 4. 3. Apparent permeability coefficient ($P_{app} \times 10^{-6}$ cm/s) of insulin across the monolayer comprising CCD18-Co, Caco-2, HT29-MTX cells and Raji B lymphocytes. Results are the mean of three replicates \pm standard deviation (n=3, mean \pm SD).

<i>In vitro</i> models	P_{app} ($\times 10^{-6}$ cm/s)	Significance level
Caco-2	2.77 ± 0.18	***
Caco-2/Raji B	5.51 ± 0.34	
CCD18-Co + Matrigel/Caco-2/HT29-MTX	7.42 ± 0.28	ns
CCD18-Co + Matrigel/Caco-2/HT29-MTX/Raji B	8.82 ± 0.98	

Comparing all the 3D models, both P_{app} values of CCD18-Co+Matrigel/Caco-2/HT29-MTX and CCD18-Co+Matrigel/Caco-2/HT29-MTX/Raji B models were significantly different from CCD18-Co+Matrigel/Caco-2 model revealing that the introduction of HT29-MTX cells was a critical factor in the permeability of insulin (Fig. 4. 28). In contrast, the P_{app} values of CCD18-Co+Matrigel/Caco-2/HT29-MTX and CCD18-Co+Matrigel/Caco-2/HT29-MTX/Raji B models were not significantly different but was observed an increase of P_{app} result of the presence of M-like cells.

Overall, the non proliferative character of CCD18-Co cells embedded in matrigel is a critical factor for the barrier formation. All the 3D models showed higher permeation compared with the respective monolayer of control. In static conditions, the presence of mucus and the less tight character between the epithelial cells contributed to the increase of insulin permeability. Arguably, there is a remote chance of insulin take advantage of M-like cells transcytotic capacity. It also remains the question whether the M-like cells formation alter the tight character of Caco-2 cell monolayer.

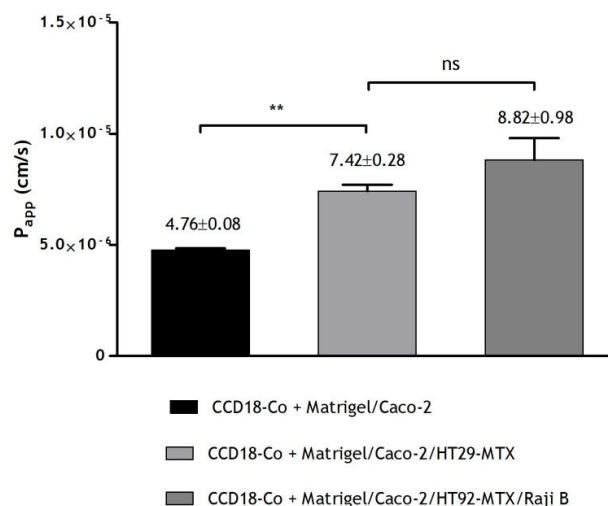


Figure 4. 28. Apparent permeability coefficient (P_{app}) of insulin through the tridimensional models comprising CCD18-Co, Caco-2 cells, HT29-MTX cells and Raji B cells. The 3D quadruple model presented the one with higher permeation to insulin. Significant differences were analysed regarding the CCD18-Co+Matrigel®/Caco-2 model. Results are the mean of three replicates and bars represent the standard deviation ($n=3$, mean \pm SD); (*) denotes a significant decrease ($p<0.05$).

Besides the low TEER values of the *in vivo* intestinal tissue, there are a plenty of other barriers that limit the absorption of the luminal content and justify the higher *ex vivo* P_{app} values. For instance, the drastic pH changes, the enzymatic activity, the presence of efflux transporters are not usually replicated in standard cell culture [56].

It is worth to mention that despite the high variability between the permeability studies of different laboratories, our data is correlated with several *ex vivo* data (Table 4. 4). When compared the P_{app} values of the 3D quadruple model with the P_{app} of *ex vivo* data, was concluded that the 3D quadruple model was efficient in predicting insulin permeability.

Table 4. 4. Apparent permeability coefficient ($P_{app} \times 10^{-6}$ cm/s) of insulin across rat ileum *ex vivo* models. Results are the mean of three replicates \pm standard deviation ($n=3$, mean \pm SD).

<i>Ex vivo</i> tissue	P_{app} ($\times 10^{-6}$ cm/s)	Reference
Ileum	0.90 ± 0.10	[170]
Ileum	0.55 ± 0.06	[171]
Jejunum	0.60 ± 0.02	[171]
Jejunum	7.35 ± 1.18	[172]

4.2.8.2. FITC-Dextran permeation

After permeability studies, the integrity of the membrane was assessed by permeating a 4 kDa small fluorescent molecule that follows the paracellular route of transport to

transverse the epithelial layer. FITC-Dextran crosses the cell layer through passive paracellular transport being a good indicative of the integrity of the membrane [173]. As observed in Fig. 4. 29, after the drug transport assays, all of the models retained the FITC-Dextran above 70%, with exception of CCD18-Co monolayer. The percentage of rejection regards the amount of molecule that the barrier retains in the apical side. This integrity studies showed that insulin had not negative repercussions in the membrane and that the tight junctions opening was reversible.

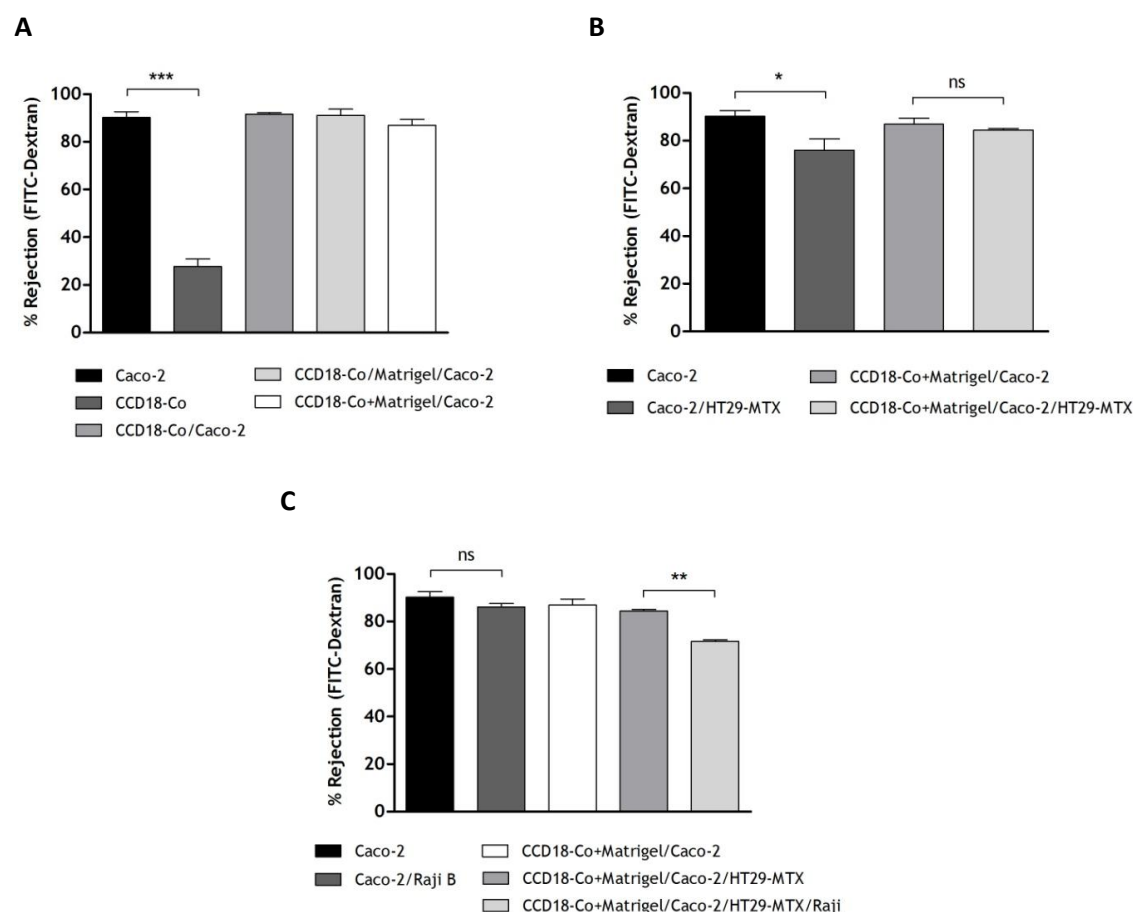


Figure 4. 29. 4 kDa FITC-Dextran permeability after 4 hours of permeability studies to assess the membrane integrity. Results are the mean of three replicates \pm standard deviation ($n=3$, mean \pm SD).

Overall, the 3D quadruple model comprising CCD18-Co intestinal myofibroblasts, Caco-2, HT29-MTX and M-like epithelial cells was morphologically and functionally characterized.

CHAPTER V

CONCLUDING REMARKS AND
FUTURE PERSPECTIVES

The herein work is motivated in satisfy the current needs of the drug development field. 'Petri-dish'-based models are limiting the potential of predict cellular responses of real organisms as they do not replicate the architecture and the mechanisms representative of a living tissue. Based on the Caco-2 gold standard model, we launched the challenge of create a third dimension that could bridge the gap between cell culture models and live intestinal tissues. In this sense, a 3D co-culture model comprising CCD18-Co intestinal myofibroblasts, Caco-2, HT29-MTX and Raji B cells was established. Beyond the fact that Caco-2 cells resembles the main features of enterocytes, this model is valued by the presence of HT29-MTX mucus producing cells in the same proportions found *in vivo* and M-like cells, an important player in the process of drug absorption owing a high transcytotic capacity. Moreover, the scenario of interdependency between stromal and epithelial cells was evaluated as mean to dissect their role in the barrier formation.

By exploring the behaviour of CCD18-Co cells in a 3D matrix through cell viability, metabolic activity, proliferation and morphological assays, 1×10^4 cells/cm² showed to be the cell density better adapted for 21 days of culture, period necessary to fully differentiate the Caco-2 cells. Added to that, CCD18-Co cells proved to be active in the modulation the surrounding Matrigel matrix through the production of fibronectin. This newly-formed matrix exerts a supportive role in the maintenance of the epithelial cells onto CCD18-Co cells. This event was the bottom line to establish a 3D model that resembles the native mucosa.

The barrier functionality is essential for purposes of absorption or permeation studies. The formation of a tightly-knit monolayer between all the cell types is determinant for the validation of results in permeability assays. Caco-2 cells form a tight polarized monolayer resulting in the highest TEER values. Even though Caco-2 and HT29-MTX cells establish tight junctions between each other, they present structural variability, resulting in a decrease of TEER values. Regarding the interactions between M-like cells and the Caco-2 and HT29-MTX cells, little is known. But the models comprising M-like cells registered a decrease in TEER values, which was probably due to the transformation of phenotype of Caco-2 cells into cells with M-phenotype. Nonetheless, this assumption requires further studies namely the identification soluble factors responsible for the phenotype induction and TJ strength.

Discriminate Caco-2 and HT29-MTX cells is difficult due to their similar morphology. Notwithstanding, HT29-MTX cells were distinctly identified by transmission electron microscopy (TEM) exhibiting the intracellular presence of mucin granules. Evidences of mucin-filled granules were found to richly cover the epithelial surface. Morphologically distinct from enterocytes, M cells display poorly organized brush border with short irregular microvilli. Even though trigger M-like cells have revealing a challenge because of the inexistence of a distinct cell marker, the lack of microvilli was still the best cue to identify these cells. Alkaline phosphatase is an excellent marker of crypt-villus differentiation, since ALP of Caco-2 cells have identical sequence and similar biochemical characteristics compared to the intestinal alkaline phosphatase (IAP). M cells exhibit down-regulation of alkaline phosphatase activity. However, there are no *in vitro* evidences of down-regulation of this

enzyme upon the phenotypic transformation. Therefore, the sensitivity of ALP assay to trigger the existence of M-like cells is doubtful and further assays are required.

The utmost importance of this work points to the sustainable results that the present model returned, supporting the successfully establishment of a barrier model reliable for the study of drug permeability. The 3D quadruple model (CCD18-Co+Matrigel/Caco-2/HT29-MTX/Raji B) was efficient in the prediction of insulin permeability when compared with several *ex vivo* models. Overall, the less tight character between the epithelial cells, the presence of mucus and M-like cells were determinant factors in the transport of insulin.

After 4 hours of permeability studies, 4 kDa FITC-Dextran molecule was retained on its majority by the barrier models, with exception of CCD18-Co monocultures, revealing that tight junctions opening was reversible and insulin did not have negative repercussions on the model integrity.

Finally, as future perspectives is imperative to understand how to control the thickness of the fibroblastic layer and quantify the amount of other ECM components. Concerning the variety of epithelial cells, it is also necessary to assess the final proportion of all the cell types in culture and understand how the presence of other epithelial cells (HT29-MTX and M-like cells) can influence Caco-2 cells differentiation. The mechanisms by which insulin cross the Caco-2/Raji B models are not fully understood. It remains to explore the TJ strength between Caco-2 and M-like cells and the contribution of the transcytotic route through M-like cells in the transport of insulin. To accomplish the undoubtful identification of M-like cells will be explored other cell markers. With this, we will be able to understand how the initial cell density of Raji B cells affects the amount of M-like cells. Finally, is necessary to test a higher number of compounds with different BCS properties and correlate these results with *in vivo* experiments.

CHAPTER VI

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